

```

acgggcgga tagagtggct taattctcga tctatcccca cgtatgcac tgaattaaca 360
aatgaactgc ttaaaaaaga cggtaagggt caagccacaa attcatttag cggagttaac 420
tattggctag ttaaaaaataa aattgaagtt ttttatccag gcccgggaca cactccagat 480
aacgtagtgg tttggttgcc tgaaaggaaa atattattcg gtggttggtt tattaaaccg 540
tacggtttag gcaatttggg tgacgcaaat atagaagctt ggccaaagtc cgccaaatta 600
ttaaagtcca aatatggtaa ggcaaaactg gttgttccaa gtcacagtga agttggagac 660
gcatcactct tgaaacttac attagagcag gcggttaaag ggttaaacga aagtaaaaaa 720
ccatcaaaac caagcaacta a 741

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<210> 1434

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1434

cacaatcaag accaagattt gcgat 25

<210> 1435

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1435

gaaagggcag ctcggttacga tagag 25

<210> 1436

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1436

cagcatcaac atttaagatc ccca 24

<210> 1437

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1437

ctccacttga ttaactgcgg aaattc 26

<210> 1438

<211> 828

<212> DNA

<213> Escherichia coli

<400> 1438
atggcaatcc gaatcttcgc gatacttttc tccatttttt ctcttgccac tttcgcgcat 60
gcgcaagaag gcacgctaga acgttctgac tggaggaagt ttttcagcga atttcaagcc 120
aaaggcacga tagttgtggc agacgaacgc caagcggatc gtgccatgtt ggtttttgat 180
cctgtgcgat cgaagaaacg ctactcgcct gcacgcacat tcaagatacc tcatacactt 240
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gccggtcgca actggatact gcgtgcaaag acgggctggg aaggccgcat ggttgggtgg 660
gtaggatggg ttgagtggcc gactggctcc gtattcttcg cactgaatat tgatacgcca 720
aacagaatgg atgatctttt caagagggag gcaatcgtgc gggcaatcct tcgctctatt 780
gaagcggttac cgcccaaccc ggcagtcaac tcggacgctg cgcgataa 828

<210> 1439
<211> 801
<212> DNA
<213> *Pseudomonas aeruginosa*

<400> 1439
atgaaaacat ttgccgcata tgtaattatc gcgtgtcttt cgagtacggc attagctggt 60
tcaattacag aaaatacgtc ttggaacaaa gagttctctg ccgaagccgt caatggtgtc 120
ttcgtgcttt gtaaaagtag cagtaaatcc tgcgtacca atgacttagc tcgtgcatca 180
aaggaatata ttccagcatc aacattttaag atccccaacg caattatcgg cctagaaact 240
ggtgtcataa agaatgagca tcaggttttc aaatgggacg gaaagccaag agccatgaag 300
caatgggaaa gagacttgac cttaagaggg gcaatacaag tttcagctgt tcccgtattt 360
caacaaatcg ccagagaagt tggcgaagta agaatgcaga aataccttaa aaaattttcc 420
tatggcaacc agaataatcag ttggtggcatt gacaaattct ggttgaagg ccagcttaga 480
atttccgcag ttaatcaagt ggagtttcta gagtctctat atttaaataa attgtcagca 540
tctaaagaaa accagctaata agtaaaagag gctttggtaa cggaggcggc acctgaatat 600
ctagtgcatt caaaaactgg tttttctggg gtgggaactg agtcaaatcc tgggtgcgca 660
tggtgggttg ggtgggttga gaaggagaca gaggtttact ttttcgcctt taacatggat 720
atagacaacg aaagtaagtt gccgctaaga aaatccattc ccaccaaact catggaaagt 780
gagggcatca ttggtggcta a 801

<210> 1440
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1440
agaccgttat cgtaaacagg gctaag 26

<210> 1441
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1441
tttttgcctc aaactttttc aggatc 26

<210> 1442

<211> 927
 <212> DNA
 <213> *Pseudomonas aeruginosa* strain RNL-1

```

<400> 1442
atgaatgtca ttataaaagc tgtagttact gcctcgacgc tactgatggt atcttttagt 60
tcattcgaaa cctcagcgca atccccactg ttaaaagagc aaattgaatc catagtcatt 120
ggaaaaaaag ccactgtagg cgttgcagtg tgggggcctg acgatctgga acctttactg 180
attaatcctt ttgaaaaatt cccaatgcaa agtgtattta aattgcattt agctatgttg 240
gtactgcatc aggttgatca gggaaagtgt gatttaaatac agaccgttat cgtaaacagg 300
gctaaggttt tacagaatac ctgggctccg ataatagaaag cgtatcaggg agacgagttt 360
agtgttccag tgcagcaact gctgcaatac tcgggtctcgc acagcgataa cgtggcctgt 420
gatttgttat ttgaactggg tgggtggacca gctgctttgc atgactatat ccagtctatg 480
ggtataaagg agaccgctgt ggtcgcaaat gaagcgcaga tgcacgccga tgatcagggt 540
cagtatcaaa actggacctc gatgaaagggt gctgcagaga tcctgaaaaa gtttgagcaa 600
aaaacacagc tgtctgaaac ctgcgaggct ttgttatgga agtggatggt cgaaaccacc 660
acaggaccag agcgggttaa aggtttgtta ccagctggta ctgtggtcgc acataaaact 720
ggtacttcgg gtatcaaagc cggaaaaact gcggccacta atgatttagg tatcattctg 780
ttgcctgatg gacggccctt gctggttgct gtttttgtga aagactcagc cgagtcaagc 840
cgaaccaatg aagctatcat tgcgcagggt gctcagactg cgtatcaatt tgaattgaaa 900
aagctttctg ccctaagccc aaattaa 927
    
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<210> 1443
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

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<400> 1443
cttctgctct gctgatgctt ggc 23
    
```

<210> 1444
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

```

<400> 1444
ggcgaccagg tattttgtaa tactgc 26
    
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<210> 1445
 <211> 927
 <212> DNA
 <213> *Salmonella typhimurium* strain JMC

```

<400> 1445
atgaatgtca tcacaaaatg tgttttcacc gcttctgctc tgctgatgct tggcttaagt 60
tcattttag tagcagccca atcccccttg ttaaaagagc agattgaaac catagtgcag 120
ggtaaaaagg ccactgtagg ttagcagtg tgggggcctg acgatctgga acctttgttg 180
ctgaatccat ttgaaaagtt tccgatgcaa agtgtgttta aactgcattt agctatgtta 240
gttctgcatc aggtcgatca ggggaaactg gatttaaatac agtctgttac tgttaatcgt 300
gctgcagtat tacaaaatac ctggtcgcca atgatgaaag atcatcaggg cgatgaatgt 360
actggttgcat tacagcagtt actgcagtat tcgggtgtcac acagcgacaa tgtggcctgc 420
gatttggttat ttgaactggg gggcgggccc caagctttgc atgcttatat ccagtcttta 480
ggcgttaaag aagctgccgt ggtagcaaat gaagcgcaaa tgcattgcgga tgatcagggt 540
caatatcaaa actggacgct gatgaaagcc gcagcacaag ttctgcaaaa gtttgaacag 600
aaaaagcagt tgtctgaaac ctctcaggcc ttgttatgga aatggatggt tgaaaccacc 660
    
```

```
acaggaccac agcgggttaaa aggcttggtta cctgctggta ctatagtggc gcataaaacc 720
ggtacttcgg gcgtcagagc aggaaaaact gcggcgacta atgatgcggg cgtcattatg 780
ttgcctgatg gacggccttt attggtggcg gtatttgtca aggattcggc tgaatcagaa 840
cgaaccaatg aagctattat tgcgcagggt gcgcaagcgg cttatcagtt tgagctgaaa 900
aaactctctg cagtgagtc ggattga 927
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<210> 1446
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1446
ggcctgygat ttgttatattg aactggt 27

<210> 1447
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1447
cgctstgggtc ctgtggtggt ttc 23

<210> 1448
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1448
gatcaggtgc artatcaaaa ctggac 26

<210> 1449
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1449
agcwggtaac aaycctttta accgct 26

<210> 1450
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:

Oligonucleotide

<400> 1450
accactggga atacacttgt aatggc 26

<210> 1451
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1451
atctacctgg tcaatcattg cttcgt 26

<210> 1452
<211> 486
<212> DNA
<213> Staphylococcus epidermidis strain BM10393

<400> 1452
atgacattat caataattgt cgctcacgat aaacaaagag tcattgggta ccaaaatcaa 60
ttaccttggc acttaccaaa tgatttaaag catattaaac aactgaccac tgggaataca 120
cttgtaatgg cacggaaaac ttttaattct ataggggaagc cattgccaaa tagacgtaac 180
gtcgtactca ctaaccaagc ttcatttcac catgaagggg tagatgttat aaactctctt 240
gatgaaatta aagagttatc tggcatgtt tttatatttg gaggacaaac gttatacgaa 300
gcaatgattg accaggtaga tgatatgtat atcacagtaa tagatggaaa gtttcaagga 360
gacacattct ttccaccata cacattcgaa aactgggaag tcgaatcttc agtagaaggt 420
caactagatg aaaaaaatac tataccgat acattcttac atttagtgcg tagaaaaggg 480
aaatag 486

<210> 1453
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1453
atcgaagaat ggagttatcg graatg 26

<210> 1454
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1454
cctaaaaytr ctggggattt cwgga 25

<210> 1455
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1455
caggtggtgg ggagatatac aaaa 24

<210> 1456
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1456
tatgtagas rcgaagtctt ggktaa 26

<210> 1457
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1457
caaaggtgaa cagctcctgt tt 22

<210> 1458
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1458
tccgttatatt tcttttaggtt ggttaaa 27

<210> 1459
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1459
aaggtgaaca gtcctgttt 20

<210> 1460
<211> 22
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1460
gatcactacg ttctcattgt ca 22

<210> 1461
<211> 474
<212> DNA
<213> Escherichia coli

<400> 1461
gtgaaactat cactaatggt agctatatcg aagaatggag ttatcgggaa tggccctgat 60
attccatgga gtgccaaagg tgaacagctc ctgtttaaag ctattaccta taaccaatgg 120
ctgttggttg gacgcaagac ttttgaatca atgggagcat tacccaaccg aaagtatgcg 180
gtcgtaacac gtccaagttt tacatctgac aatgagaacg tagtgatctt tccatcaatt 240
aaagatgctt taaccaacct aaagaaaata acggatcatg tcattgtttc aggtgggtggg 300
gagatatata aaagcctgat cgatcaagta gatacactac atatatctac aatagacatc 360
gagccggaag gtgatgttta ctttcctgaa atccccagca attttaggcc agtttttacc 420
caagacttcg cctctaacat aaattatagt taccaaatct ggcaaaaaggg ttaa 474

<210> 1462
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1462
gcactcccy aataggaaata cgc 23

<210> 1463
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1463
agtgttgctc aaaaacaact tcg 23

<210> 1464
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1464
acgttygaat ctatgggmgc act 23

<210> 1465
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1465
gtcgataagt ggagcgtaga ggc 23

<210> 1466
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1466
aagcattgac ctacaatcag tgt 23

<210> 1467
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1467
aatacaacta cattgtcatc atttgat 27

<210> 1468
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1468
cgttacccgc tcaggttgga catcaa 26

<210> 1469
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1469
catccccctc tggctcgatg tcg 23

<210> 1470
<211> 474
<212> DNA
<213> Escherichia coli

<400> 1470

ttgaaagtat	cattgatagc	tgcgaaacga	aaaaacggcg	tgattgggtg	cggtccagac	60
ataccgtggg	ccgcgaaagg	ggagcagcta	ctttttaaag	cattgaccta	caatcagtgt	120
cttctgggtg	gtcgcaagac	gtttgaatct	atgggcgcac	tccccaatag	gaaatacgcg	180
gtcgttaccc	gctcagggtg	gacatcaaat	gatgacaatg	tagttgtatt	tcagtcaatc	240
gaagaggcca	tggacaggct	agctgaattc	accgggtcacg	ttatagtgtc	tggtggcgga	300
gaaatttacc	gagaaacatt	acccatggcc	tctacgctcc	acttatcgac	gacgacatc	360
gagccagagg	gggatgtttt	cttcccagag	attccaaata	ccttcgaagt	tgtttttgag	420
caacacttta	cttcaaacat	taactattgc	tatcaaattt	ggaaaaaggg	ttaa	474

<210> 1471
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1471	
gataatgaca acgtaatagt attccc	26

<210> 1472
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1472	
gctcaatatc aatcgctcgat ata	23

<210> 1473
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1473	
ttaaagcctt gacgtacaac cagtgg	26

<210> 1474
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1474	
tgggcaatgt ttctctgtaa atctcc	26

<210> 1475
 <211> 474
 <212> DNA
 <213> Escherichia coli

```

<400> 1475
gtgaaagtat cattaatggc tgcaaaagcg aaaaacggag tgattgggtg cgggtccacac 60
ataccctggg ccgcgaaagg agagcagcta ctcttttaaag ccttgacgta caaccagtgg 120
cttttggtgg gccgcaagac gttcgaatct atgggagcac tccctaatag gaaatacgcg 180
gtcgttactc gctcagcctg gacggccgat aatgacaacg taatagtatt cccgtcgatc 240
gaagaggcca tgtacgggct ggctgaactc accgatcacg ttatagtgtc tgggtggcggg 300
gagattttaca gagaaacatt gcccatggcc tctacgctcc atatatcgac gattgatatt 360
gagccggaag gagatgtttt ctttcggaat attcccaata ctttcgaagt tgtttttgag 420
caacacttta gctcaaacat taactattgc tatcaaattt ggcaaaaggg ttaa 474

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<210> 1476

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1476

```
ggcgagcagc tcctattcaa ag 22
```

<210> 1477

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1477

```
taggtaagct aatgccgatt caaca 25
```

<210> 1478

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1478

```
gagaatggag taattggctc tggatt 26
```

<210> 1479

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1479

```
gcgaaataca caacatcagg gtcatt 25
```

<210> 1480

<211> 474

<212> DNA

<213> Proteus mirabilis strain J120

```
<400> 1480
atgaaaatat ctcttatggc agctggtttcc gagaatggag taattggctc tggattggat 60
ataccttggc atgtacaagg cgagcagctc ctattcaaag ccatgactta caatcaatgg 120
cttctagttg gtcgtaaaac cttcgactca atgggtaaac ttccgaatag aaaatatgca 180
gtggttactc gttctaaaat tatctcgaat gaccctgatg ttgtgtatgt cgcaagtgtt 240
gaatcggcat tagcttacct aaacaatgcg acagcacata tctttgtttc tgggtgggtggt 300
gaaatatata aagctttaat cgatcaagca gatgttatcc atctttcagt gattcacaag 360
catatctctg gcgatgtgtt ttttcctcca gttccacagg gcttcaagca aacatttgag 420
caaagtttca gttcaaatat tgattacacg taccaaattt gggcaaaggg ctaa 474
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<210> 1481

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

```
<400> 1481
rttacagatc atktatatgt ctct 24
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<210> 1482

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

```
<400> 1482
taatttatat tagacawaaa aaactg 26
```

<210> 1483

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

```
<400> 1483
carygtcaga aaatggcgta atc 23
```

<210> 1484

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

```
<400> 1484
tkcaaagcrw tttctattga aggaaa 26
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<210> 1485

<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1485
aaaatggcgt aatcggtaat ggc 23

<210> 1486
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1486
catttgagct tgaaattcct ttcctc 26

<210> 1487
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1487
aatcgaaaat atgcagtagt gtcgag 26

<210> 1488
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1488
agactattgt agatttgacc gccca 24

<210> 1489
<211> 474
<212> DNA
<213> Escherichia coli strain VA292

<400> 1489
ttgaaaattt cattgatttc tgcaacgtca gaaaatggcg taatcggtaa tggccctgat 60
atcccatggt cagcaaaaagg tgagcagtta ctcttttaaag cgctcacata taatcagtgg 120
ctccttggtg gaaggaaaac atttgactct atgggtgttc ttccaaatcg aaaatatgca 180
gtagtgtcga ggaaaggaat ttcaagctca aatgaaaatg tattagtctt tccttcaata 240
gaaatcgctt tgcaagaact atcgaaaatt acagatcatt tatatgtctc tgggtggcgg 300
caaattctaca atagtcttat tgaaaaagca gatataattc atttgtctac tgttcacggt 360
gaggttgaag gtgatatcaa ttttcctaaa attccagaga atttcaattt ggtttttgag 420
cagttttttt tgtctaatat aaattacaca tatcagattt ggaaaaaagg ctaa 474

<210> 1490
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1490
gacctatgag agcttgcccg tcaaa 25

<210> 1491
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1491
tcgccttcgt acagtcgctt acaaaa 26

<210> 1492
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1492
catttttagct gccaccgcca atggtt 26

<210> 1493
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1493
gcgtcgctga cgttgttcac gaaga 25

<210> 1494
<211> 510
<212> DNA
<213> Escherichia coli strain BL26A

<400> 1494
atgatcgagc ttcatgccat tttagctgcc accgccaatg gttgcattgg gaaggacaac 60
gcacttcctt ggccaccact aaaaggcgat ctggccagat tcaaaaaatt gaccatgggg 120
aaggtggtca ttatggggcg caagacctat gagagcttgc ccgtcaaatt agaagggtcgc 180
acctgcatcg ttatgacgcg ccaagcgctg gagcttcctg gtgttcgtga cgctaacggc 240
gctatcttcg tgaacaacgt cagcgacgcc atgcggttcg ctcaagaaga gagcgtgggc 300
gatgtggcct acgtcattgg tggcgctgag atattcaagc gacttgcctt gatgatcacg 360
cagattgaat tgacctttgt taagcgactg tacgaaggcg acacctacgt tgatctggcc 420
gaaatgggtca aagactacga gcagaatggc atggaagaac atgaccttca cacttacttc 480

acttaccgta aaaaggagct tacagaatga 510

<210> 1495
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1495 24
 tctctaaaca tgattgtcgc tgtc

<210> 1496
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1496 24
 cagtgaggca aaagtttttc tacc

<210> 1497
 <211> 25
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1497 25
 cggacgactt catgtggtag tcagt

<210> 1498
 <211> 26
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1498 26
 tttgttttca gtaatggtcg ggacct

<210> 1499
 <211> 534
 <212> DNA
 <213> Escherichia coli

<400> 1499
 atggcttctc taaacatgat tgctcgtgtc aataagacag gaggtatcgg atttgaaaat 60
 cagattccgt ggcatgaacc agaagattta aaacacttca aagctgttac aatgaactca 120
 gttttgatta tgggtagaaa aacttttgcc tctactgcta aagtgtgtcc cggacgactt 180
 catgtggtag tcagtaaaac agtaccaccc acccagaaca ctgatcaagt tgtgtatgta 240
 agtacatacc agatcgcagt aagaactgca agcttggtgg ttgacaaaacc agagtatttct 300

caaatttttg taattggtgg gaagagtgcg tacgagaact tagctgccta cgtggacaaa 360
ctctacttaa ctagagtaca gctcaacaca caacaagaca ctgaactgga tttatcccta 420
ttcaagtcac ggaaactcgt atctgaggtc ccgaccatta ctgaaaacaa aacaaaactt 480
attttccaaa tttggattaa ccctaaccct attagtgagg aaccacatg ttag 534

<210> 1500
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1500
atcgggttat tggcaatggt ccta 24

<210> 1501
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1501
gcggtagtta gcttggcgtg agatt 25

<210> 1502
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1502
gcgggcggag ctgagatata ca 22

<210> 1503
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1503
aacggagtgg gtgtacggaa ttacag 26

<210> 1504
<211> 498
<212> DNA
<213> Escherichia coli strain TKS84

<400> 1504
atgaactcgg aatcagtagc catttatctc gttgctgcga tgggagccaa tcgggttatt 60
ggcaatggtc ctaatatccc ctggaaaatt ccgggtgagc agaagatttt tcgcagactc 120

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actgagggaa aagtcgttgt catggggcga aagacctttg agtctatcgg caagcctcta 180
ccgaaccgtc acacattggt aatctcacgc caagctaact accgcgccac tggctgcgta 240
gttgtttcaa cgctgtcgca cgctatcgct ttggcatccg aactcggcaa tgaactctac 300
gtcgcgggcg gagctgagat atacactctg gcactacctc acgcccacgg cgtgtttcta 360
tctgaggtac atcaaacctt cgaggggtgac gccttcttcc caatgctcaa cgaaacagaa 420
ttcgagcttg tctcaaccga aaccattcaa gctgtaattc cgtacacca ctccgtttat 480
gcgcgtcgaa acggctaa

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<210> 1505
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1505
 atttttcgca ggctcaccga gagc 24

<210> 1506
 <211> 27
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1506
 cggatgagac aacctcgaat tctgctg 27

<210> 1507
 <211> 498
 <212> DNA
 <213> Escherichia coli strain RA33.2

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<400> 1507
atgaaccgga aatcgggtccg catttatctg gtcgctgcca tgggtgcca tcgggttatt 60
ggcaatggtc ccgatatccc ctggaaaatc ccagggtgagc agaagatttt tcgcaggctc 120
accgagagca aagtgggtcgt tatgggcccgc aagacatttg agtccatagg caagccctta 180
ccaaaccgcc acacagtggg gctctcgcgc caagctgggtt atagcgctcc tggttgtgca 240
gttggtttcaa cgctgtcaca cgtatcgcca tcgacagccg aacacggcaa agaactctac 300
gtagcgcgcg gagccgaggt atatgcgctg gcgctaccgc atgccaacgg cgtctttcta 360
tctgaggtac atcaaacctt tgaggggtgac gccttcttcc cagtgcctta cgcagcagaa 420
ttcgaggttg tctcatccga aaccattcaa ggcacaatca cgtacacgca ctccgtctat 480
gcgcgtcgta acggctaa

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<210> 1508
 <211> 27
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1508
 agaattgtatt ggtatttcca tctatcg 27

<210> 1509

<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1509
caatgtcgat tgttgaaata tgtaaa 26

<210> 1510
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1510
tggagtgcc aaggggaaca at 22

<210> 1511
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1511
cagacacaat cacatgatcc gttatcg 27

<210> 1512
<211> 474
<212> DNA
<213> Escherichia coli strain UI14

<400> 1512
gtgaaactat cactaatggc agcaatttcg aagaatggag ttatcggaaa tggcccagat 60
attccatgga gtgccaaagg ggaacaatta ctcttcaaag cgattaccta taatcagtgg 120
cttttggtag gccgaaagac tttcgagtca atggggggctt tacccaaccg aaaatatgcc 180
gttgtaactc gttcaagctt cacttccagt gatgagaatg tattggtatt tccatctatc 240
gatgaagcgc taaatcatct gaagacgata acggatcatg tgattgtgtc tgggtggtggt 300
gaaatataca aaagcctgat cgataaagtt gatactttac atatttcaac aatcgacatt 360
gagccagaag gtgatgtcta ttttccagaa atccccagta gttttaggcc agtttttagc 420
caagacttcg tgtctaacat aaattatagt taccaaactt ggcaaaaggg ttaa 474

<210> 1513
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1513
ttcaagctca aatgaaaacg tcc 23

<210> 1514
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1514
gaaattctca ggcattatag ggaat 25

<210> 1515
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1515
gtggtcagta aaaggtgagc aac 23

<210> 1516
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1516
tcctttcaaag cattttctat tgaagg 26

<210> 1517
<211> 474
<212> DNA
<213> Escherichia coli strain EC107

<400> 1517
ttgaaaatat cattgatttc tgcagtgtca gaaaatggcg taatcggtag tggctcctgat 60
atcccgtggc cagtaaaagg tgagcaacta ctcttttaaag cgctcacata taatcaatgg 120
ctccttgctcg gaagaaaaac atttgactct atgggtgttc ttccaaatcg caaatatgca 180
gtagtgtcaa agaacggaat ttcaagctca aatgaaaacg tcctagtttt tccttcaata 240
gaaaatgctt tgaaagagct atcaaaaagtt acagatcatg tatatgtctc tggcgggggt 300
caaatctata atagccttat tgaaaaagca gatataattc atttgtctac tgttcacggt 360
gaagtcgaag gtgatataca attccctata atgcctgaga atttcaattt ggtttttgaa 420
cagtttttta tgtctaatat aaattataca taccagattt ggaaaaaagg ctaa 474

<210> 1518
<211> 125
<212> DNA
<213> Acinetobacter lwoffii strain CDCF 3697

<400> 1518
ctatgtctca aggcggtgca acataactcta tggaatttgc taaatatgct gaaactccac 60
gtaacgtggc tgaaggcatc atttctaaat ttcagtctgg cggtaaaaaa ggtgacgacg 120
agtaa 125

<210> 1519
 <211> 93
 <212> DNA
 <213> *Acinetobacter lwoffii* strain CDCF 3697

<400> 1519
 tctttcgatt actataagcc caaactaatt catagttaaa aaccaagtgc tcatgcagtg 60
 atcctgcatg agtagtttaa aaaggaagat ctc 93

<210> 1520
 <211> 1106
 <212> DNA
 <213> *Acinetobacter lwoffii* strain CDCF 3697

<400> 1520
 atggctaagg ctaagtttga acgtaataag ccacacgtta acgtgggcac aatcgggtcac 60
 gttgaccatg gtaaaacaac tttaacagct gcaattgcaa ctgtatgtgc gaagaaattc 120
 ggtggcgaag cgaaagacta cgctgcaatt gactctgcac cagaagaaaa agcacgtggt 180
 attacaatta atacttcaca cgtagaatac gattctccaa ctcgtaacta cgcacacgta 240
 gactgcccgg gccacgccga ttatgtttaa aacatgatta ctggtgctgc tcagatggac 300
 ggcgcgatcc ttgtatgtgc tgcgactgat ggtccaatgc cacagactcg tgaacacatc 360
 cttctttctc gtcagggttg tgtaccttac attcttgtat tccttaacaa gtgtgacctt 420
 gttgatgatg aagaacttct tgagctagtg gaaatggaag ttctgtgaact tctttctact 480
 tatgacttcc caggatgatg cactccagtt atccgtggtt cagctcttct tgcacttaac 540
 ggtgacgctg gtcagtatgg cgaagaagca gttgttgccg ttgttgacgc acttgacact 600
 tacattccag agccagtacg tgcaatcgac caagcattct taatgccaat cgaagacgta 660
 ttctctatct ctggtcgtgg tacagtagta actggccgtg tagaaactgg tattgtgaaa 720
 gtagggcgaat cagttgaaat cgttggtatc cgtgatactc aagtaactac agttactggc 780
 gtacaaatgt tccgtaaatt gcttgacgaa ggtcgtgccc gcgagaactg tgggtgttctt 840
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 atcaagccac aactaaatt cgatgcagaa gtatacgtac tttctaaaga agaaggtggt 960
 cgtcacactc cattccttaa cggttaccgt ccacagttct acttccgtac aactgacgta 1020
 actggcgcga tcaaattaca agatggcgtt gaaatgggta tgcctggtga caacgtagaa 1080
 atgtcagtag aattaatcca cccaat 1106

<210> 1521
 <211> 100
 <212> DNA
 <213> *Haemophilus influenzae* ATCC 9006

<400> 1521
 acaaaactcaa ggtcgtgcat cttactcaat ggaaccgtta aaatatgctg aagctccaac 60
 aagtgttgcg gctgcagtaa ttgaagcgcg taaaaaataa 100

<210> 1522
 <211> 64
 <212> DNA
 <213> *Haemophilus influenzae* ATCC 9006

<400> 1522
 tttttgtaaa ccagcgggtgt aaaatatgat tgttttatac cgcacttctt aggaaacatt 60
 agaa 64

<210> 1523
 <211> 1098
 <212> DNA
 <213> *Haemophilus influenzae* ATCC 9006

<400> 1523
 atgtctaaag aaaaatttga acgtacaaaa ccgcacgtaa acgtgggtac aatcggccac 60
 gttgaccacg gtaaaacaac tttaacagca gcaattacaa ccgtattagc aaaacactac 120
 ggtgggtgcag cgcgtgcatt tgaccaaatc gataacgcgc cagaagaaaa agcgcgtggt 180

attaccatca	acacttcaca	tgttgaatac	gatacaccaa	ctcgccacta	tgcacacgta	240
gactgtccag	gacacgccga	ctatgttaaa	aacatgatta	ccggtgcggc	gcaaatggat	300
ggtgctat	tagtagtagc	agcaacagat	ggctctatgc	cacaaactcg	tgaacatata	360
ttattaggtc	gccaagtagg	tgttccatac	atcatcgat	tcttaaacia	atgagacatg	420
gtagatgatg	aagagttatt	agaattagta	gaaatggaag	tgcgtgaact	tctatctcaa	480
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ggcgtagcag	aatgggaaga	aaaaatcctt	gaattagctg	gtcacttaga	tacttacatc	600
ccagaaccag	aacgtgcat	tgaccaaccg	ttccttcttc	caattgaaga	cgtattctca	660
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gatgaagtag	aaatcgctcg	tatcaaagat	acagcgaaaa	ctactgtaac	aggtggtgaa	780
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ccacacactg	atgttgaatc	agaagtatac	gtattatcaa	aagatgaagg	tggtcgctcat	960
actccattct	tcaaagggtta	ccgtccacaa	ttctatttcc	gtacaacaga	cgtactgggt	1020
acaattgaat	taccagaagg	cgtggaaatg	gtaatgccag	gcgataacat	caagatgaca	1080
gtaagcttaa	tccacca					1098

<210> 1524
 <211> 77
 <212> DNA
 <213> Proteus mirabilis ATCC 25933

<400> 1524	
caatggagtt	cttgaagtac aacgaagcgc ctagcaacgt cgctcaggct attatcgaag 60
ctcgtaaagc	gaaataa 77

<210> 1525
 <211> 67
 <212> DNA
 <213> Proteus mirabilis ATCC 25933

<400> 1525	
gaccccttcg	agttcaattt agtttacgct ccctctgtga gagggagcga tattaaggaa 60
tatagtc	67

<210> 1526
 <211> 1112
 <212> DNA
 <213> Proteus mirabilis ATCC 25933

<400> 1526	
gtgtctaaag	aaaaatttga acgttcaaaa ccgcaagttta acgttggtac tatcggccac 60
ggtgaccacg	gtaaaacaac tctgactgct gcaatcacta cagttttagc taaaacttac 120
ggtggtgctg	ctcgtgcatt cgaccaaatac gataatgcac cagaagaaaa agcgcgtggg 180
atcaccatct	ctacttcaca cgtagaatac gatactccaa ctcgccacta cgcacacgta 240
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atctcaggcc	gtggtacagt agttactggg cgtgtagagc gtggtatcat caaagtaggt 720
gatgaagttg	agattgttgg tatcaaagaa accaccaaaa caacttgtag tggcgttgag 780
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ccacacaaca	aatttgaatc agaagtttat attctgagca aagatgaagg tggtcgctcac 960
acaccattct	tcaaaggcta ccgtccacag ttctacttcc gtacaactga cgtactgggt 1020
actatcgaat	taccagaagg cgtagaaatg gtaatgccag gcgacaacgt gaacatgatc 1080
ggtgaactga	tccaccaat cgcaatggac ga 1112

<210> 1527

<211> 800
<212> DNA
<213> *Campylobacter curvus* ATCC 35224

<400> 1527
atcaacgaag ctatcgaggt ttattttgag gttgagggca agaaaaatag attgatcctg 60
gaggtcgcgg ctcaacttgg tgataaccgc gtcagaacga tcgctatgga tatgagttag 120
gggcttactc gcgggcttga agctaccgct cttgggtgcgc ctattagtgt gccggttggc 180
gagaaggttt tgggaagaat ttttaacgct gtcggcgatc tcatcgacga gggcgagggc 240
gtaaattttg ataaacattg gtctatccac cgcgatccgc caccatttga agaacaaagc 300
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gagctcatcc acaatgtcgc cttcaaacac agcggatact ctgtatttgc aggtgttggc 480
gagaggacgc gcgaaggaaa cgacctttat cacgagatga aagaaagtaa cgttttggat 540
aaagtcgcct tgtgctacgg acagatgaac gagccgccag gggcgagaaa tcgtatcgca 600
ctgactggtc taacgatggc tgagtatttc cgcgatgaga tgggacttga tgtgcttatg 660
tttatcgaca acatcttccg cttctctcaa tctggtgcag agatgtcggc actcctcggg 720
cgtatcccat cagccgttgg ttaccagccg acgctggcaa gcgagatggg taaatttcaa 780
gaaaggatca catcgactaa 800

<210> 1528
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1528
aacttgagcg attttcggat accctg 26

<210> 1529
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1529
ttgccgatga aataaccgcc gact 24

<210> 1530
<211> 1035
<212> DNA
<213> *Escherichia coli*

<400> 1530
atgcgattgg tttggaaatg tggggcgatt caggcatccc ggttatctga atgggtcaac 60
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ggctcagtgct tgatctggct gaaatcatat ctccgcgaat caggaagaaa actgcagtta 180
gtcggaatcg ccttacccaa caccctgaac ccaagggacg acctagcgca attggccgaa 240
attatccagc tcatcgatca cctcatgaaa ccgcacgttg atatggtgac tcacttggtg 300
gcgtccattg atggccagtc ggcggttatt tcatcggcaa aatgggggga gctagaaacg 360
gctcggcagg agaaagctat ctcaggggta accagattga agctccgctt ggcgtcgctt 420
gccccgtcc tgaaaaaaca cgtcaacagc gatttgttcc gaaaagcctc tgatcgaata 480
gagtcgatag agtatacgtt ggaaaccttg cgtataatga aaactttctt cgatggtacc 540
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gttcgagcga atccggatgt gaagataatt ctgctggcgc acaacaatca tctacaaaaa 660
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cgtgaggata gtatggaaca gtatgtcatc gacgcctgtg gtacggagaa ttcattgtctg 900
acattgacag atgcccccat ggaagcaaag cgaatgcggt ctcaaagcgc ctctgtagaa 960
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ctggttgccc tataag                                     1035
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<210> 1531
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1531
tctttttgtt acgacatacg ctttt 25

<210> 1532
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1532
agtgttctt tatccgctgt tcta 24

<210> 1533
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1533
cagcggataa agaagcacta cacatt 26

<210> 1534
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1534
cctcctgaaa taaagcccga cat 23

<210> 1535
<211> 1260
<212> DNA
<213> Escherichia coli

<220>
<221> misc_feature

<222> (41)..(41)
<223> n represents any nucleotide

<220>
<221> misc_feature
<222> (47)..(47)
<223> n represents any nucleotide

<220>
<221> misc_feature
<222> (93)..(93)
<223> n represents any nucleotide

<220>
<221> misc_feature
<222> (1170)..(1170)
<223> n represents any nucleotide

<400> 1535
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gataattacg atgattttta gccattaaga aanataattg gagatacccg agttgtagca 120
ttaggtgaaa attctcattt cataaaaagaa ttctttttgt tacgacatac gcttttgctg 180
ttttttatcg aagatctagg ttttactacg tttgcttttg aatttggttt tgctgagggt 240
caaatcatca ataactggat acatggacaa ggaactgacg atgaaatagg cagattctta 300
aaacacttct attatccaga agagctcaaa accacatttc tatggctaag ggagtacaat 360
aaagcagcaa aagaaaaaat cacatttctt ggcattgata taccagaaa tggagggttca 420
tacttaccaa atatggagat agtgcagac ttttttagaa cagcggataa agaagcacta 480
cacattatcg atgatgcatt taatattgca aaaaagattg attacttctc cacatcacag 540
gcagccttaa atttacatga gctaacagat tctgagaaat gccgtttaac tagccaatta 600
gctcgagtaa aagttcgcct tgaagctatg gctccaattc acattgaaaa atatgggatt 660
gataaatatg agacaattct gcattatgcc aacggtatga tatacttgga ctataacatt 720
caagctatgt cgggctttat ttcaggaggc ggaatgcagg gcgatatggg tgcaaaaagac 780
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ggttttcgag ttgataactt ccaactgcag gaaccaaattg aaggttctgt cgagaaagct 1080
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caatccatcc cgaacatgat tcgatttgan tctatttaca tgaaagcaga actcgagaaa 1200
gctttcgatg gaatatttca aattgaaaag tcatctgtat ctgaggtcgt ttatgaataa 1260

<210> 1536
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1536
agatgtatta actggaaaac aacaa

25

<210> 1537
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1537

ctttgtaatt agtttctgaa aacca

25

<210> 1538
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1538
ttagaagata taggatacaa aatagaag

28

<210> 1539
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1539
gaatgaaaaa gaagttgagc tt

22

<210> 1540
<211> 486
<212> DNA
<213> Staphylococcus haemolyticus

<400> 1540
atgaaaaata ataatgtaac agaaaaagaa ttattttata ttttagattt atttgaacac 60
atgaaagtaa cttattgggt agatgggtggc tggggggtag atgtattaac tggaaaacaa 120
caaagagaac acagagatat agatatagat tttgacgctc aacacactca aaaagttata 180
caaaaattag aagatatagg atacaaaata gaagttcatt ggatgccttc acgtatggaa 240
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gaaactaatt acaaagatcg aaaaatacca tgtatttcaa aagaagctca acttcttttt 420
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acataa 486

<210> 1541
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1541
tgataatctt atacgtgggg aattt

25

<210> 1542
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:

Oligonucleotide

<400> 1542
ataattttct aattgccctg tttcat 26

<210> 1543
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1543
gggcaattag aaaattatatt atcaga 26

<210> 1544
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1544
ttttactcat gtttagccaa ttatca 26

<210> 1545
<211> 804
<212> DNA
<213> Enterococcus faecium

<400> 1545
atgttaaaac aaaaagaatt aattgcaaac gttaagaatc ttactgagtc agatgaacga 60
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cgattagata aggtagaatt atttgaagcc tataaaaaat ctttgctatt agttatggat 720
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ttgttgaatt acattagtga atag 804

<210> 1546
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1546
caagaaggaa tggctgtact ac 22

<210> 1547
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1547
taattcccaa ataaccctaa taataga

27

<210> 1548
<211> 1218
<212> DNA
<213> Streptococcus pyogenes

<400> 1548
atggaaaaat acaacaattg gaaacttaag ttttatacaa tatgggcagg gcaagcagta 60
tcattaatca ctagtgccat cttgcaaatg gcgattatatt ttaccttac agaaaaaact 120
ggatccgcga tgggtcttgc tatggcttca ctattaggtt ttttacccta tgcggtcttt 180
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ggctatagtc agtctttgca gtctataagc tatattgtta gtccggcggt tgcagcactc 480
ttatactccg tttgggaact aaatgctatt attgccatcg atgtattggg tgctgtgatt 540
gcatctatta cggtagcaat tgtacgtatt cctaagctgg gtgatcgcg gcaaagtgtg 600
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aaattagatt taaaataa 1218

<210> 1549
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1549
gcttattatt aggaagatta gggggc

26

<210> 1550
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1550
tagcaagtga catgatactt ccga

24

<210> 1551
<211> 1218
<212> DNA
<213> Streptococcus pneumoniae

<400> 1551
atggaaaaat acaacaattg gaaacgaaaa ttttatgcaa tatgggcagg gcaagcagta 60
tcattaatca ctagtgccat cctgcaaatg gcgattattt tttaccttac agaaaaaaca 120
ggatctgcga tgggtctgtc tatggcttca ttagtaggtt tttacccta tgcgattttg 180
ggacctgcca ttggtgtgct agtggatcgt catgatagga agaagataat gattgggtgcc 240
gattttaatta tcgcagcagc tgggtgcagt cttgctattg ttgcattctg tatggagcta 300
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ctatcaggta ttttaattat tggcattgct atagtttgcc aaatgataac tgagggttaga 1200
aaattagatt taaaataa 1218

<210> 1552
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1552
ggcaagcagt atcattaatc acta 24

<210> 1553
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1553
caatgctacg gataaacaat actatc 26

<210> 1554
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1554

agaaaattaa gcctgaatat ttaggac

27

<210> 1555

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1555

tagtaaaaac caatgattta caccg

25

<210> 1556

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1556

actgtacgca cttgcagccc gacat

25

<210> 1557

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1557

gaacggcagg cgattcttga gcat

24

<210> 1558

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1558

gtggtggtgc atggcgatct ct

22

<210> 1559

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1559

gccgcagcga ggtactcttc gtta

24

<210> 1560
 <211> 906
 <212> DNA
 <213> *Escherichia coli*

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<400> 1560
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gccaccgtcg acgatggacg tcgggtgggtg ctgcgcaccc cgcgccgagc cgaggtaagc 180
gcgaaggctc aaccagaggg gcgggtgctg gcaatgctca agaatcgctt gccgttcgctg 240
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agcgaggccc gcgttgatga ccctgccatc gacatggccg cgcaccttat ggtctttggt 720
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cggtcgcgcc accacatcgc ggagcgctt gcgttcgggg cggtcaccta cgcactcttc 840
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gaatga                                           906
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<210> 1561
 <211> 1048
 <212> DNA
 <213> *Candida albicans* ATCC 18804

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<400> 1561
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gccaaacttt tggattaygg ttctattgat agagctccag aagaaagagc tagaggatc 120
actattttcc ctgcccacgt tgaatacgaa accaagaaca gacactatgc ccacgttgat 180
tgtccaggac acgctgatta tatcaaaaat atgattactg gtgccgctca aatggatgg 240
gctatcattg ttgttctgct cactgatggg caaatgcctc aaaccagaga acatttggt 300
ttggccagac aagttgggtg tcaagacttg gttgtgtttg tcaacaaaag cgatactatt 360
gatgaccctg aaatgttggg attagtcgaa atggaaatga gagaattggt atccacctac 420
ggttttgatg gtgacaacac tccagttatt atgggactcg ctttaattgg tttggaagac 480
aagaaaccag aaatttggtg ggaagctatc ttgaaattgt tagatgctgt cgatgaacac 540
attccaactc catcaagaga cttggaacaa ccatttttgt taccagttga agacgtgttc 600
tccatctccg gtagaggaac tgttgtcact ggttagagtg aaagaggtgt tttgaagaag 660
ggtgaagaaa tcgaaattgt tgggtggttt gacaaacctt acaagactac tgttaccggt 720
attgaaatgt tcaaaaaaga attagactct gctatggctg gtgacaactg tgggtgtttg 780
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cgttccactc catttggtga aggttacaag cctcaatgct tcttcagaac taacgatgtc 960
actaccacat tttcattccc agaaggagaa ggtgttgacc attctcaaat gatcatgcca 1020
ggtgacaaca ttgaaatggg ttggtgaat                                           1048
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<210> 1562
 <211> 1074
 <212> DNA
 <213> *Candida dubliniensis* strain NCPF 3949

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<400> 1562
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tgccaacttt ttgattacg gttccattga tagagctcca gaagaaagag ccagaggtat 120
cactattttc actgcccacg ttgaatacga aaccaagaac agacactatg cccacgttga 180
ttgtccagga cacgctgatt atatcaaaaa catgattact ggtgctgctc aaatggatgg 240
tgctatcatt gttgttctg ctactgacgg tcaaatgcca caaaccagag aacatttatt 300
gttggaaga caagttgggt ttcaagactt ggttgccttt gtcaacaaag ttgatactat 360
tgatgacct gagatgttgg aattagtcga aatggaaatg agagaattgt tgtccacct 420
cggttttgat ggtgacaaca ctctgttat tatgggatct gctttaatgg ccttggagg 480
caaaaaacca gaaattggta aggaagctat tttgagattg ttagatgctg tcgatgaaca 540
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cattccaact ccatcaagag acttggaaaca accattttttg ttgccagttg aagacgtggt 600
ctccatctct ggtagaggaa ctgttgtcac cggtagagtt gaaagaggtg tcttgaagaa 660
gggtgaagaa atcgaaattg ttggtgggtt tgacaaacca tacaagacca ctgttactgg 720
tattgaaatg ttcaaaaagg aattagattc tgctatagct ggtgacaact gtggtggttt 780
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tgctacttct cacaagaaat ttttagcatc ttgttatatt ttgacttcag aagaagggtgg 900
tcgttccact ccatttggag aagggttacia gcctcaatgt ttcttcagaa ctaatgacgt 960
cactaccaca ttttcattcc cagaaggaga aggtgttgac cactcccaa tggtcacgac 1020
aggtgataac attgaaatgg ttggtgaatt gatcaaata tgtccattgg aagt 1074
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<210> 1563
<211> 1033
<212> DNA
<213> *Candida famata* ATCC 62894

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<400> 1563
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gctaacttct tggactacgg ttctatcgat aaagctccag aagaaagagc cagagggtatt 120
actatttctg ctgcccattg tgaatacgaa actgacaaga gacactatgc ccatgttgat 180
tgtccagggt acgcagatta tatcaagaat atgattactg gtgctgctca aatggatggg 240
gccattattg ttgttgctgc ttccgatggg caaatgcctc aaaccagaga acatttggtt 300
ttggccagac aagttggtgt tcaacacttg gttgttttcg tcaacaagggt cgacaccatt 360
gacgatccag aaatgttgga attggttgaa atggaaatga gagatttggt aactacttac 420
ggttttgatg gtgataacac cccagttatc atgggatctg ctttgtgtgc tttggaatcc 480
agagaaccag aaattggtca aaaagccatt gaaaattgt tagatgccgt cgatgaatac 540
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accggtacca tgaccttccc agaaggtgcc gaccaatctg ccatggtcat gccagggtgac 1020
aacgttgaaa tgc 1033
```

<210> 1564
<211> 1056
<212> DNA
<213> *Candida glabrata* ATCC 66032

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<400> 1564
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gctgatttct tggactactc ttccattgac aaagctccag aggagagagc ccgtgggtatc 120
actatctcta ctgcccattg cgagtaacag accgccaaga gacattactc ccacgtcgac 180
tgtccagggt acgccgacta catcaagaac atgattactg gtgctgcca aatggacggg 240
gctatcatcg ttgtcgccgc caccgatggg caaatgccac aaactagaga gcatttgctg 300
ttggccagac aagtcgggtg tcaacgtatc gttgtctttg tcaacaagggt ggacaccatc 360
gatgaccctg aaatgttgga attagtgga atggaaatga gagaattggt gaacgaatac 420
ggttttgacg gtgacaatgc ccctatcatt atgggttccg ctttgtgtgc cctagaagggt 480
cgtcaacctg aaattggtga gcaagctatc atgaaactat tggacgctgt tgatgaatac 540
attccaaccc cagaaagaga cttgaacaag ccattcttga tgccgttgta agacattctc 600
tccatctctg gtagaggtac cgtcgctact ggacgtgtcg aaagaggtaa cttgaagaag 660
ggtgaagaag ttgaaattgt tggtcacaac actaccccat tgaagaccac cgttactggg 720
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agacattctg gtttcgggtg aaactacaga cctcagatgt ttatcagaac cgcagatgtc 960
actgttggtg tgaagttccc agaactctgt gaagaccact ctatgcaagt tatgccagggt 1020
gacaacgctg aaatgggtcg tgaactagtc caccaca 1056
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<210> 1565
<211> 1061
<212> DNA

<213> *Candida guilliermondii* ATCC 6260

<400> 1565

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accatttcca	ctgcccattg	tgagtaccaa	actgataaga	gacattatgc	ccacgttgac	180
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gccattattg	ttgttgctgc	caactgacgg	caaatgcctc	agaccagaga	gcacttggtg	300
ttggccagac	aagttgggtg	gcaacacttg	gtagtttttg	tgaacaagg	ggacaccatt	360
gacgatcccc	agatgtttga	attggctcgag	atggaaatga	gagaattggt	gagtcagtac	420
ggtttcgatg	gtgacaacac	cccagttatc	atgggatctg	ctttgtgtgc	tttggaaagt	480
aagcagccag	aaattgggtg	gcaagccatt	gaaaaattgt	tggacgctgt	cgatgagcac	540
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tccatttctg	gtagaggaac	tgtggttact	ggtagagtcg	aaagagggtc	gttgaagaag	660
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cgtcactctc	cttttgggtg	gaactacaag	cctcaattgt	tcacagaac	tactgacgtt	960
accggtactt	taagattccc	agccggcgag	ggtgtcgacc	actcgcaaat	ggttatgcc	1020
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<210> 1566

<211> 1073

<212> DNA

<213> *Candida haemulonii* ATCC 22991

<400> 1566

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<210> 1567

<211> 1062

<212> DNA

<213> *Candida kefyr* ATCC 28838

<400> 1567

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ccaacccag	cccgtgactt	ggaaaavcca	ttcttgatgc	ctggtgaaga	tatcttctcc	600
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gaagaaatcg	aaattgttgg	tcacaacacc	actcctttca	agactactgt	tactgggtatt	720
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gtdgtcttga	agttcccaga	atctgttgaa	gaccattcca	tgcaagtcac	gccaggtgac	1020
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<210> 1568
 <211> 1062
 <212> DNA
 <213> *Candida lusitaniae* ATCC 66035

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aggtcacgcc	gattacatca	agaacatgat	cacgggtgcc	gctcaaattg	acggtgccat	240
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ctctccattt	ggcgagaact	acaagcctca	attgttcctt	agaactaccg	atgtcactgg	960
tactttgaga	ttcccagcag	gtgaggacgt	tgaccactcc	gctatggttt	ctccaggtga	1020
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<210> 1569
 <211> 990
 <212> DNA
 <213> *Candida sphaerica* ATCC 2504

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<210> 1570
 <211> 1184
 <212> DNA
 <213> *Candida tropicalis* ATCC 750

<400> 1570

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ctaacgatgt	taccacttcc	ttctctttcc	cagaagggtga	aggtgttgac	cactcccaaa	1140
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<210> 1571

<211> 1071

<212> DNA

<213> *Candida viswanathii* ATCC 28269

<400> 1571

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cactatctcc	actgcccacg	ttgaatacga	gactgataag	agacactatg	cccacgttga	180
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caccaccact	ttcactttcc	cagaagggtga	aggtgtcgac	cactcccaaa	tggttatgccc	1020
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<210> 1572

<211> 817

<212> DNA

<213> *Alcaligenes faecalis* subsp. *faecalis* ATCC 8750

<400> 1572

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gccacacact	gacttcgacg	ccgaggtgta	cattctgtcc	aaagaagaag	gtgggtcgta	660
cactcctttc	ttcaagggtc	accgtcctca	gttctacttc	cgtacaactg	acgtgaccgg	720

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agtgtccctg atcgctccta tcgccatgga agaaggt 817

<210> 1573
<211> 796
<212> DNA
<213> *Prevotella buccalis* ATCC 35310

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<210> 1574
<211> 820
<212> DNA
<213> *Succinivibrio dextrinosolvens* ATCC 19716

<400> 1574
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atcgatctga aagagggcgt agagatggta atgccaggtg ataacaccga catgaccgta 780
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<210> 1575
<211> 803
<212> DNA
<213> *Tetragenococcus halophilus* ATCC 33315

<400> 1575
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aaaattctca gctgaagttt atgtattaac aaaagaagaa ggcggacgtc atactccatt 660
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attgccagaa ggtactgaaa tggttatgcc aggtgataac gtagcaatgg aagttgaatt 780
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<210> 1576

<211> 805

<212> DNA

<213> *Campylobacter jejuni* subsp. *jejuni* ATCC 33292

<400> 1576

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<210> 1577

<211> 809

<212> DNA

<213> *Campylobacter rectus* ATCC 33238

<400> 1577

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<210> 1578

<211> 1671

<212> DNA

<213> *Enterococcus casseliflavus* ATCC 25788

<400> 1578

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gcgtgttgta	ttctgtaaca	aaatggacaa	aattggtgca	gacttcttat	actctgtatc	180
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tgtaccagaa	gattacttag	gtgatgttat	gggtcagcta	actgctcgtc	gtggacgcgt	1620
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<210> 1579

<211> 1662

<212> DNA

<213> *Enterococcus gallinarum* ATCC 49573

<400> 1579

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<210> 1580

<211> 1669

<212> DNA

<213> *Streptococcus mitis* ATCC 49456

<400> 1580

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<210> 1581
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1581
aattggggac tacacctatt atgatg 26

<210> 1582
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1582
ggcaaatcag tcagttcagg agt 23

<210> 1583
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1583
cgattggcaa caatacactc ctg 23

<210> 1584
<211> 26
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1584

tcacctatatt ttacgcctgg taggac

26

<210> 1585

<211> 645

<212> DNA

<213> Enterococcus faecium

<400> 1585

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gccaaccacg	taatgaaagg	tatttcgact	tatccattta	atatttttagg	tggtcgattgg	300
caacaatata	ctcctgaact	gactgatttg	ccgttgaaag	gtgatactgt	agtcggaaat	360
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aatccaattc	aactcatcgg	accaagattt	gaaccggaag	ttattcaagc	attagaaaaat	540
ctggcatggg	ggaataaaga	tattgaatgg	ataactgcta	atgttcctaa	actaatgcaa	600
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<210> 1586

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1586

attcccacaa tcttttttat caataa

26

<210> 1587

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1587

cattgttcag attcggtaaa gttc

24

<210> 1588

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1588

gtttttgaag ttaaatagtg ttctt

25

<210> 1589
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1589
cttccatttg tactttccct a

21

<210> 1590
<211> 1920
<212> DNA
<213> Unknown Organism

<220>
<223> Description of Unknown Organism: Cloning vector
pFW16

<400> 1590
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gcaaaagatg gcgtacaagc acaaactcgt atattatttc atgcacttag gaaaatgggg 360
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<210> 1591
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

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<400> 1591
atgaggtaat agaacggatt
20

<210> 1592
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1592
cagtattttca gtaagcgtaa a
21

<210> 1593
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
<221> misc_feature
<222> (29)..(29)
<223> n represents a modified base

<220>
<221> modified_base
<222> (29)..(29)
<223> i

<400> 1593
ccgagcgcatt taccggatac ttggctgcnc gctcgg
36

<210> 1594
<211> 1032
<212> DNA
<213> Enterococcus faecium strain N97-330

<400> 1594
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cttaggaggt aa
1032

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<210> 1595
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1595
aaataatgct ccatcaattt gctga 25

<210> 1596
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1596
atagtcgaaa aagccatcca caag 24

<210> 1597
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1597
gatgaatttg cgaaaatata tgga 24

<210> 1598
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1598
cagccaattt ctaccctttt cac 23

<210> 1599
<211> 604
<212> DNA
<213> Enterococcus faecalis strain BM4405

<400> 1599
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caattgggtcg ttggagaatg tgatcaaatt agtcttggtg atggcttttt cgactatgaa 420
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aaagaagaca tccagataaa agcaaaaaaa ctatacagac tattaggggtg caaaggatta 540
gcgagaatcg acttttttctt aacggatgac ggagaaatgt tattaaatga gatcaacacc 600
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<210> 1600

<211> 805

<212> DNA

<213> *Campylobacter jejuni* subsp. *doylei* ATCC 49349

<400> 1600

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gcttatgttt attgataata tcttttagatt ttcacaatca ggttctgaaa tgtcagcact 720
tttaggaaga attccatcag ctgtgggtta tcaaccaacc ctagcaagtg aaatgggtaa 780
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<210> 1601

<211> 826

<212> DNA

<213> *Enterococcus sulfureus* ATCC 49903

<400> 1601

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atcgactgat ggattgcaac gtgggatgga agttgtcgat atgggagaaat ctatttctgt 180
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tgctttgtta ggtcgtatgc catcagctgt gggatatcaa ccaacattag ctaccgaaat 780
ggggcaattg caagagcgga tcacgtcaac gaaaaaagga tcgatc 826

<210> 1602

<211> 833

<212> DNA

<213> *Enterococcus solitarius* ATCC 49428

<400> 1602

tgatacttta ccagatatta ataatgcatt agtagtatat aaaaaggacg aggacaagac 60
acgcgttggtc ttagaagcca ccttggaact tggagatggc atgattcgtg caatctctat 120
gggatcgact gatggcttgc aacggggaat ggaagttgtg gacacacaag cacctatttc 180
tgttccagta ggaaatgaaa ccttaggacg tgtttttaat gtcttaggag aaacgattga 240
taaacaggca ccgttttctg aagatgccaa aaaaagtggg attcataaaa aagcaccgcg 300
ttttgatgaa ttaagtacca gttctgaaat attggaaacc gggattaaag taatcgattt 360
gctagctcct tatttacgag gtggtaaaat tggattattt ggcggtgctg gcgtgggtaa 420
aacggtatta attcaagaat taattcataa cgttgcccaa gaacatgggg gaatttctgt 480
ttttacgggt gtcggagagc gtactcgtga aggaaatgac ctatattatg aaatgcagga 540
ttcaggcggt attgaaaaaa cggctatggg atttggacaa atgaacgaac cccctgggtg 600
acgtatgcgt gtagcggtta ctggtttgac acttgctgag tacttccgtg atgtacaagg 660

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tcaagacgta ttattattta tagataatat tttccgcttt actcaagcag gaacagaagt 720
atctgcttta ttaggacgga tgccgtctgc cgttgggttac caaccaactc tagcaacgga 780
aatgggacag ttgcaagaac gaatcacatc gacagataaa ggatcaatta cct 833
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<210> 1603

<211> 806

<212> DNA

<213> *Campylobacter sputorum* subsp. *sputorum* ATCC 35980

<400> 1603

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gacttattct tgaagtagct ggacatcttg gcgataatag agcaagaacc attgctatgg 120
atatgagtga tggtttacaa agaggattag aagttacggc tcttggtgct cctataacag 180
ttcctggttg agataaagtt ttaggtagaa tgtttaatgt tgtaggtgac ttaatagatg 240
aaggtgaagt aacagatttt gataaaagat gggctatcca tagagatcct ccttcgtttg 300
aagatcaaag tacaaaaagt gaaatttttg aaacaggtat aaaagtagtt gatcttcttg 360
caccttatgc aaaaggtggt aaagttggct tatttggtgg tgctggcggt ggaaaaacag 420
ttatcataat ggagcttata cataatgttg catttaaaca cagcggttat tcaatttttg 480
ccggtggttg agagagaaca agagaggaa atgatcttta taatgagatg aaagagtctg 540
gtgttttggg taaagttgcc ttatgttatg gacaaatgaa tgaaccacca ggagcaagaa 600
accgtatagc attaacaggt cttacaatgg ctgaatatgt ccgtgatgaa atgggggcttg 660
atgtgttgat gtttatagat aatattttta gattttctca atcaggttct gaaatgtcag 720
cgctgcttgg tagaattcca tctgctgttg gttatcaacc aacattagca agtgagatgg 780
gaaaacttca agaaagaatt acttcc 806
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<210> 1604

<211> 738

<212> DNA

<213> *Enterococcus pseudoavium* ATCC 49372

<400> 1604

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ggtgttatcc gcacaatcgc tatggaatct acagatggat tgcaacgggg gatggaagtt 60
gtcgataccg gcaaaccaat ctctgttcct gtaggtaaag aaacattagg tcgtgtgttt 120
aacgtattag gtgaaacgat cgacaaagaa gcaccttttc cagaagatgt agaaaagagc 180
ggtattcaca aaaaggcccc cgcttttgaa gaccttagca ccagtaatga gatttttagaa 240
actgggatca aggttatcga cttattagcc cttacttaa aaggtggtta agttggacta 300
ttcggtggtg ccggtgttgg taaaaccgtc ttaattcaag aactgattca taatatcgcc 360
caagaacacg gtgggatttc tgtctttacc ggggttgggg aacggactcg tgaagggaac 420
gacctttatt atgaaatgaa agaatccggc gttattgaaa aaacagcgat ggtcttcgga 480
caaatgaatg agccaccagg tgcgcggatg cgcggtgcct tgactggttt gacattagct 540
gaatatttcc gtgatgaaga aggtcaagat gtgttgctat ttatcgataa cattttccgc 600
ttcacacaag ccggtacaga agtttcggcg ctattaggtc ggatgccatc tgccggttgg 660
tatcaaccaa ccttggcaac agaaatgggt caattacaag aacgaatcac ttcaacgaaa 720
aaaggctcaa ttacatcg 738
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<210> 1605

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (12)..(12)

<223> n represents a modified base

<220>

<221> misc_feature

<222> (18)..(18)

<223> n represents a modified base

<220>
 <221> modified_base
 <222> (12)..(12)
 <223> i

<220>
 <221> modified_base
 <222> (18)..(18)
 <223> i

<400> 1605
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20

<210> 1606
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
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 <222> (9)..(9)
 <223> n represents a modified base

<220>
 <221> misc_feature
 <222> (12)..(12)
 <223> n represents a modified base

<220>
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 <223> n represents a modified base

<220>
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 <222> (9)..(9)
 <223> i

<220>
 <221> modified_base
 <222> (12)..(12)
 <223> i

<220>
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 <222> (15)..(15)
 <223> i

<400> 1606
 ccraacatna yncnacttt ttc

23

<210> 1607
 <211> 336
 <212> DNA
 <213> Klebsiella ornithinolytica ATCC 31898

<400> 1607
 ctggattatg cgatgtcggc cattgttggc cgtgcgctgc cggatgtccg agatggcctg 60
 aaaccggtac accgtcgcgt actttacgcc atgaacgtat tgggcaatga ctggaacaaa 120
 gcctataaaa aatccgcccc tgctgttggc gacgtaatcg gtaaatacca ccctcatggt 180

gataccgccc tttatgacac cattgtacgt atggcacagc cattctcctt gcggttatatg 240
ctggtcgatg gccagggtaa cttcggttct gtcgatggcg actccgccgc agcgatgcgt 300
tatacggaaa tccgtatgtc gaaaatcgcc cacgag 336

<210> 1608
<211> 341
<212> DNA
<213> Klebsiella oxytoca ATCC 13182

<400> 1608
ctatctggat tatgcatgt cggtcattgt tggccgtgcg ctgccggatg tccgagatgg 60
cctgaagccg gtacaccgtc gcgtactata cgccatgaac gtattgggca atgactggaa 120
caaagcctat aaaaaatctg cccgtgtcgt ggggtgacgtc atcggtaa at accaccctca 180
tggtgatact gccgtatacg acaccattgt acgtatggcg cagccattct cctgcggtta 240
catgctggta gatggccagg gtaactttgg ttcggtcgac ggcgactccg ccgcagcgat 300
gcggtatacg gaaatccgta tgtcgaagat cgcccatgaa c 341

<210> 1609
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1609
gccctgatcc aaatagcata ta 22

<210> 1610
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1610
cctggcataa cagtaacatt ctg 23

<210> 1611
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1611
tgggaaaaag caactccatc tc 22

<210> 1612
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1612
acaactgaat tcgcagcaac aat

23

<210> 1613
<211> 639
<212> DNA
<213> Staphylococcus aureus

<400> 1613
atgaaatatg gccctgatcc aaatagcata tatccacatg aagaaataaa aagtgtttgt 60
tttattaaaa atacaattac caatccaaat attatagttg gagattatac ttactattcc 120
gatgttaacg gagctgaaaa atttgaagaa catgtgacac atcattatga atttaggggt 180
gataaacttg taattggcaa gttttgtgca atagctgaag gtatagaatt tattatgaat 240
ggagcaaacc atagaatgaa ttcaataaca acttatcctt ttaatataat gggaaatggg 300
tgggaaaaag caactccatc tcttgaagat ttaccattta agggagatac tgttggttga 360
aatgatgtgt ggattgggtca gaatgttact gttatgccag gaattcaaatt aggagatgga 420
gcaattgttg ctgcgaattc agttgttaca aaagatgtac caccatatcg tattattggg 480
ggaaatccga gtagaattat aaagaaaagg tttgaagatg aattgataga ttattttattg 540
caaataaaat ggtgggattg gtcagcacia aaaatatttt ctaatcttga aacactttgt 600
agctctgatt tagagaaaat aaaatctatt cgagattag 639

<210> 1614
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1614
ccaatccaga agaaatatac cc

22

<210> 1615
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1615
attagtttat cccaatcaa ttca

24

<210> 1616
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1616
ataatgaatg gggctaataca tcgtat

26

<210> 1617
<211> 24
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1617

gccacaact gaataaggat caac

24

<210> 1618

<211> 639

<212> DNA

<213> Staphylococcus cohnii

<400> 1618

atgaaatggc	aaaatcagca	aggccccaat	ccagaagaaa	tataccctat	agaaggtaat	60
aaacatgttc	aattttattaa	accatctata	acaaagccca	atatttttagt	tgggggaatat	120
tcatattacg	atagtaaaga	tggatgaatct	tttgaaagcc	aagttcttta	tcactatgaa	180
ttgattgggg	ataaactaat	attaggggaag	ttttgttcta	ttggaccggg	aacgacattt	240
ataatgaatg	gggctaataca	tcgtatggat	ggttcaacat	ttccattcaa	tcttttcgga	300
aatggttggg	agaagcatac	ccctacattg	gaagaccttc	cttataaggg	taacacggaa	360
attgggaacg	atgtttggat	tggacgagat	gtgacaatta	tgcccgggtgt	aaaaatagga	420
aacggggcta	ttattgcagc	aaaatcggtt	gtgacaaaga	acgttgatcc	ttattcagtt	480
gttggcggtg	atccttcacg	attaattaag	ataaggtttt	ccaaggaaaa	aatcgcagca	540
ttactaaaag	taagggtggtg	ggacctagag	atagagacga	taaatgaaaa	tattgattgc	600
atcctgaatg	gtgatataaa	aaagggttaa	agaagtttag			639

<210> 1619

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1619

aaggcaaaat aaaaggagca aagc

24

<210> 1620

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1620

tgtacccgag acatcttcac cac

23

<210> 1621

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1621

aattgaagga cgggtattgt ggaaag

26

<210> 1622
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1622
cgattttgac agatggcgat aatgaa

26

<210> 1623
<211> 1569
<212> DNA
<213> Staphylococcus aureus

<400> 1623
atgaaaataa tgttagaggg acttaatata aaacattatg ttcaagatcg tttattgttg 60
aacataaatc gcctaaagat ttatcagaat gatcgtattg gtttaattgg taaaaatgga 120
agtggaaaaa caacgttact tcacatatta tataaaaaaa ttgtgcctga agaagggtatt 180
gtaaaacaat tttcacattg tgaacttatt cctcaattga agctcataga atcaactaaa 240
agtgggtggtg aagtaacacg aaactatatt cggcaagcgc ttgataaaaa tccagaactg 300
ctattagcag atgaaccaac aactaactta gataataact atatagaaaa attagaacag 360
gatttaaaaa attggcatgg agcattttatt atagtttcac atgatcgcgc ttttttagat 420
aacttgtgtgta ctactatatg ggaaattgac gagggaagaa taactgaata taagggggaat 480
tatagtaact atgttgaaca aaaagaatta gaaagacatc gagaagaatt agaatatgaa 540
aaatatgaaa aagaaaagaa acgattggaa aaagctataa atataaaaga acagaaagct 600
caacgagcaa ctaaaaaacc gaaaaactta agtttatctg aaggcaaaat aaaaggagca 660
aagccatact ttgcaggtaa gcaaaagaag ttacgaaaaa ctgtaaaatc tctagaaacc 720
agactagaaa aacttgaaag cgtcgaaaag agaaacgaac ttctccact taaaatggat 780
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acaattgaag gacgggtatt gtggaaagca aaaagtttta gtattcgcgg aggagacaag 900
atggcaatta tcggatctaa tggtagagga aagacaacgt ttattaaaaa aattgtgcat 960
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gatacattag aattagataa gagcatttta gaaaatgttc aatcttcttc acaacaaaat 1080
gaaactctta ttcgaactat tctagctaga atgcattttt ttagagatga tgtttataaa 1140
ccaataagtg tcttaagtgg tggagagcga gttaaagtag cactaactaa agtattctta 1200
cgtgaagtta atacgttggg actagatgaa ccaacaaact ttcttgatat ggaagctata 1260
gagggcgtttg aatctttgtt aaaggaatat aatggcagta taatctttgt atctcacgat 1320
cgtaaattta tcgaaaaagt agccactcga ataatgacaa ttgataataa agaaataaaa 1380
atattttgatg gcacatatga acaattttaa caagctgaaa agccaacaag gaatattaaa 1440
gaagataaaa aacttttact tgagacaaaa attacagaag tactcagtcg attgagtatt 1500
gaaccttcgg aagaattaga acaagagttt caaaacttaa taaatgaaaa aagaaatttg 1560
gataaataa 1569

<210> 1624
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1624
ttctttaatg ctcgtagatg aaccta

26

<210> 1625
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1625
ttttcgtatt cttcttggtg ctttc 25

<210> 1626
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1626
aggaatgatt aagccccctt caaaaa 26

<210> 1627
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1627
ttacattgcg accatgaaat tgctct 26

<210> 1628
<211> 1658
<212> DNA
<213> Staphylococcus aureus

<400> 1628
atgcttaaaa tcgacatgaa gaatgtaaaa aaatattatg cagataaatt aattttaaat 60
ataaaagaac taaagattta tagtggggat aaaataggta ttgtaggtaa gaatggagtt 120
ggcaaaacaa cactttttaa aataataaaa ggactaatag agattgacga aggaaatata 180
attataagtg aaaaaacaac tattaaatat atctctcaat tagaagaacc acatagtaag 240
ataattgatg gaaaatatgc ttcaatatat caagttgaaa ataagtggaa tgacaatatg 300
agtgggtggtg aaaaaactag atttaaacta gcagagggat ttcaagatca atgttcttta 360
atgctcgtag atgaacctac aagtaattta gatatcgaag gaatagagtt gataacaaat 420
acttttaaag agtaccgtga tacttttttg gtagtatctc atgatagaat ttttttagat 480
caagtttgta caaaaatttt tgaaattgaa aatggatata ttagagaatt catcggtaat 540
tatacaaact atatagagca aaaagaaatg cttctacgaa agcaacaaga agaatacgaa 600
aagtataatt ctaaaagaaa gcaattggag caagctataa agctaaaaga gaataaggcg 660
caaggaatga ttaagcccc ttcaaaaaca atgggaacat ctgaatctag aatatggaaa 720
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atagataaat taaatcatgt agaaaaaata aaagagcttc cttctattaa aatggattta 840
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gtagaatcag taataatatc accatcagtt aaaattggat acgtcagtc aaacttagat 1080
gttctacaat ctcataaatc tatcttagaa aatgttatgt ctacctccat tcaagatgaa 1140
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gcattagaag aattgttaat tacctatgaa ggtgttggtg tatttgcttc ccatgataaa 1380
aaattttatac aaaacctagc tgaacaattg ttaataatag aaaataataa agtgaaaaaa 1440
ttcgaaggaa catatataga atatttaaaa attaaagata aaccaaattt aaatacaaat 1500
gaaaaagaac tcaaaagaaa aaagatgata ctagaaatgc aaatttcac attattaagt 1560

aaaatctcaa tggaagaaaa tgaagaaaaa aacaaagaat tagatgaaaa gtacaaattg 1620
aaattaaaaag aattgaaaag cctaaataaa aatattta 1658

<210> 1629
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1629 26
aaggggaaag tttggattac acaaca

<210> 1630
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1630 24
gaaccacagg gcattatcag aacc

<210> 1631
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1631 21
cgacgatgct ttatggtttg t

<210> 1632
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1632 26
gttaatttgc ctatcttgtc acactc

<210> 1633
<211> 900
<212> DNA
<213> Staphylococcus aureus

<400> 1633
atggaattta aattacaaga attaaatctt actaaccaag atacaggacc atatggtata 60
accgtttcag ataaggggaa agtttggatt acacaacata aagcaaata gataagttgc 120
atcaatttag atggaaaaat tacagagtac ccactaccga caccagatgc aaaagtcacg 180
tgtttaacta tatcctcaga tggggaagtt tggtttactg agaatgcagc aaacaaaata 240

gggaggatta	caaaaaaagg	gattattaag	gaatatacat	tgcctaacc	agattcagca	300
ccctacggta	ttacagaagg	accaaatgga	gatatatggt	ttacagaaat	gaatggcaac	360
cgtattggac	gtattacgga	cgacggtaaa	attcgtgaat	acgagctgcc	taataaagga	420
tcttaccctt	cttttatcac	tttgggttct	gataatgccc	tgtggttcac	agaaaatcaa	480
aataatgcta	ttggtagaat	tacagaaagt	ggggatatta	cagagttaa	aattcctaca	540
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attatcggta	ataagatagg	gcgaataact	cctctggggg	aaattaccga	attcaaaatt	660
ccaacgccaa	acgctcgacc	tcatgcaatt	actgctggag	caggaattga	tttatggttt	720
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ccaattcaaa	tcaaaagtgg	tgaaccacat	ggcatttgtt	tcgatgggtga	aacaatttgg	840
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<210> 1634
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1634
ttaacttgtc tattccccgat tcagg 25

<210> 1635
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1635
gctgtggcaa tggatattct gta 23

<210> 1636
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1636
ttcctacccc tgatgctaaa gtga 24

<210> 1637
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1637
caaagtgcgt tatccgaacc taa 23

<210> 1638

<211> 527
<212> DNA
<213> *Aspergillus fumigatus* strain WSA-172

<400> 1638
ggtggggaag actgtcttca tccaggaatt gattgtgagt cgttccacat gtcacctag 60
ttttcgctcg atctttttcac taacgcaaac catgtagaac aacattgcca aggcccacgg 120
tggttactcc gtcttctactg gtgttgggtga gcgtactcgt gagggtaacg atctgtacca 180
cgaaatgcag gagactgggtg tcatttcagct cgaggggtgaa tccaagggtcg cactgggtgtt 240
cggacagatg aacgagcccc ccggtgcccc tgcccgtgtc gcccttaccg gtctgaccat 300
tgccgagtac ttccgtgacg aggaggggtca ggacgtgctg ctcttcattg acaacatttt 360
ccgttttcacc caggccggtt ctgaggtgtc tgcccttctc ggtcgtatcc cctctgccgt 420
cggttaccag cccaccctgg ccgtcgacat ggggtggtatg caggagcgta tcaccaccac 480
caagaagggt tctattacct ccgtccargc cgtctacgtc cccgcga 527

<210> 1639
<211> 452
<212> DNA
<213> *Aspergillus fumigatus* ATCC 64746

<400> 1639
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ttctgtgggtg ttgggtgaacg tactcgtgaa ggtaacgatt tataccacga aatgatggaa 120
actgggtgtca ttaaacttga aggtgactcc aagtgtgctc ttgtattcgg tcaaatgaac 180
gaacctcctg gtgctcgtgc ccgtgttgct ttaactgggt taaccattgc tgaatacttc 240
cgtgatgaag aaggtcaaga tgtgttactt ttcattgata acattttccg tttcactcaa 300
gctgggttctg aagtatctgc ccttttaggt cgtattccat ctgctgtagg ttaccaaccc 360
actttatcta ctgatatggg tggatgcaa gaacgtatta ctactaccaa gaatgggttc 420
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<210> 1640
<211> 783
<212> DNA
<213> *Bacillus mycoides* ATCC 6462

<400> 1640
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cacttggtcg tgtattcaac gtattaggtg atgcaattga cttagatggt gaacttcctg 240
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caggtttaac aatggctgag catttcctgt atgagcaagg acaagacgta cttctgttca 660
tcgataacat cttccgtttc acgcaagcgg gttctgaagt atctgccctt cttggctcgt 720
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gta 783

<210> 1641
<211> 823
<212> DNA
<213> *Bacillus mycoides* NRRL NRS-319

<400> 1641
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gatggacttg ttcgtggcac agaagtagaa gatactggta aagcaatctc tgtaccagtt 180
ggtagtgcaa cacttggtcg tgtattcaac gtattaggtg atgcaattga cttagatggt 240
gatgttcctg cggatgtacg tcgtgatcca attcaccgtc aagcacctgc attcgaagaa 300
ctatctacta aagtagaaat tcttgaaact ggtattaaag tagtagactt acttgctcct 360

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gtaggtgagc	gtactcgtga	gggtaacgac	ttataccatg	aaatgagcga	ttctggcgta	540
attaagaaaa	ctgcgatggg	attcggacaa	atgaatgagc	cacctggagc	acgtcaacgt	600
gttgcattaa	cagggtttaac	aatgggtgaa	catttcctgt	atgagcaagg	acaagacgta	660
ctattgttca	tcgataacat	cttcggtttc	acgcaagcag	gttctgaagt	atctgccctt	720
cttggtcgta	tgccatctgc	ggtaggttac	caaccaacac	ttgcaacaga	aatgggtcaa	780
ttacaagagc	gtattacatc	tacaaataaa	ggatctatca	cgt		823

<210> 1642
 <211> 829
 <212> DNA
 <213> *Bacillus mycoides* NRRL BD-15

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aaagcaatct	ctgtaccagt	tggtgatgta	acacttggtc	gtgtattcaa	cgtattaggt	240
gatgcaattg	acttagatgg	tgaagttcct	cgggatgtac	gtcgtgatcc	aattcaccgt	300
caagcacctg	cattcgaaga	attatctact	aaagtagaaa	ttcttgaaac	tggtattaaa	360
gtagtagact	tacttgctcc	ttacattaag	gggtggaaga	ttggtctatt	cggtgggtgcc	420
gggtgtaggta	aaacagtatt	aattcaggaa	ttaattaaca	acatcgca	agaacacggt	480
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gaaatgagcg	attctggcgt	aattaagaaa	actgcatgg	tattcgga	aatgaacgag	600
ccacctggag	cacgtcaacg	tgttgcat	acaggtttaa	caatggctga	gcatttcctg	660
gatgagcaag	gacaagacgt	actactgttc	atcgataaca	tcttcctgtt	cacgcaagca	720
ggttctgaag	tatctgccct	tcttggtcgt	atgccatctg	cggtaggtta	ccagccaaca	780
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<210> 1643
 <211> 823
 <212> DNA
 <213> *Bacillus pseudomycoides* NRRL BD-10

<400> 1643						
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gttatctacg	aaagtagaaa	ttcttgaaac	tggtattaaa	gtagtagact	tacttgctcc	360
ttacattaaa	gggtgtaaaa	tcggtctatt	cggtgggtgcc	gggtgtaggta	aaacagtatt	420
aatccaggaa	ttaattaaca	acatcgca	agagcacggt	ggtatttctg	tattcgctgg	480
tgtaggtgag	cgtactcgtg	aaggtaatga	cttataccac	gaaatgagcg	attctggcgt	540
aatcaagaaa	acagcgatgg	tattcgga	aatgaacgag	ccacctgggtg	cacgtcaacg	600
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tcttggtcgt	atgccatctg	cggtaggtta	ccaaccaact	cttgcaacag	aaatgggtca	780
attacaagag	cgtattacat	ctacaaata	aggatctatc	acg		823

<210> 1644
 <211> 708
 <212> DNA
 <213> *Bacillus pseudomycoides* NRRL B-617

<400> 1644						
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tacattaaag	gtggtaaaaat	cggtctatct	ggtgggtgccg	gtgtaggtaa	aacagtatta	420
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gtaggtgagc	gtactcgtga	aggtaacgac	ttataccatg	aaatgagcga	ttctggcgta	540
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gttgcattaa	caggtttaac	aatggctgaa	catttccgtg	atgagcaagg	acaagacgta	660
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<210> 1645

<211> 778

<212> DNA

<213> *Budvicia aquatica* ATCC 35567

<400> 1645

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gaaagtactt	gatttagaac	accctatcga	agttcctgtc	ggtgtggcaa	ctctggggccg	180
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ggaaacgggc	atcaagggtta	tcgacctgat	ttgtccggtt	gccaaaggcg	gtaaagtgtg	360
tctgtttggt	ggtgccggcg	taggtaaaaac	ggtaaactatg	atggagctga	ttcgtaatat	420
tgcgactgag	cactcaggtt	actctgtatt	tgcggcgctt	ggtgagcgta	ctcgtgaggg	480
taatgacttc	taccacgaaa	tgacagaatc	taacgtatta	gacaaagtat	ctctgggttta	540
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ttatacctta	gccggtacag	aagtatcagc	actgttaggt	cgtatgccat	cagcggtagg	720
ttaccaacca	acgctggcgg	aagagatggg	tacactgcaa	gaacgtatca	cytcaacc	778

<210> 1646

<211> 806

<212> DNA

<213> *Buttiauxella agrestis* ATCC 33320

<400> 1646

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tctagctccc	aggatctgct	ggaaaccggc	atcaaagtaa	tggacctgat	ttgcccgttc	360
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cgtatgccat	ctgcggtagg	ttaccagcca	actctggcag	aagagatggg	tgttttgcag	780
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<210> 1647

<211> 1122

<212> DNA

<213> *Candida norvegica* ATCC 36586

<400> 1647

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ggtaacgatt	tataccgtga	aatgaaagaa	accgggtgtca	ttaacttggg	aggtgactct	420
aaagtcgctt	tagtcttcgg	tcaaataaac	gaacctccag	gtgctagagc	ccgtgttgcc	480
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ttgaaagaaa ccgttgcttc attcagagac gtttttagctg gt 1122

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<210> 1648
 <211> 813
 <212> DNA
 <213> *Streptococcus pneumoniae* ATCC 700677

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aaccaagctg tcggaaccaa tcgtgactgg ggatcaacta tgaaaccaat cacagactat 180
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<210> 1649
 <211> 813
 <212> DNA
 <213> *Campylobacter lari* ATCC 43675

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<210> 1650
 <211> 570
 <212> DNA
 <213> *Coccidioides immitis* strain WSA-222

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ttcactgggt tcggtgagcg taccctgtgag ggtaacgatt tgtaccatga aatgcaagag 180
acctgtgtca ttcaactcga cggagagtcc aaggctcgctc ttgtcttcgg tcaaatgaac 240

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gagccccctg	gtgcccgtgc	ccgtgttgcc	cttaccggtt	tgaccattgc	tgaatacttc	300
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aggaggctaa	tcgcttttct	agtgtctctc	tttattgaca	acattttccg	tttcaactca	420
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actctcgccg	tcgacatggg	tgttatgcag	gaacgtatca	ccaccaccac	caagggatcc	540
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<210> 1651

<211> 560

<212> DNA

<213> *Emmonsia parva* ATCC 10784

<400> 1651

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gactcgaagc	tccactcaca	catatattag	tgctcctctt	catcgacaac	attttccgct	420
tcacccaggc	aggttccgaa	gtgtccgccc	tgctcgcccg	tatccccctc	gccgtcggtt	480
accagcccac	cctcgctgtc	gacatgggta	tgatgcagga	acgtatcacc	accaccacca	540
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<210> 1652

<211> 780

<212> DNA

<213> *Erwinia amylovora* ATCC 14976

<400> 1652

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<210> 1653

<211> 545

<212> DNA

<213> *Fonsecaea pedrosoi* ATCC 18831

<400> 1653

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ccgtc						545

<210> 1654

<211> 564
 <212> DNA
 <213> *Fusarium moniliforme* strain WSA-213

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gctgagtact tcagagacga ggagggtcag gacgtgctgc ttttcattga caacattttc 420
cgattcactc aggcgcgttc cgagggtgtc gcccttctcg gtcgtatccc ctctgccgtc 480
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accaagggtt ccattacctc agtc                                     564
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<210> 1655
 <211> 776
 <212> DNA
 <213> *Klebsiella oxytoca* ATCC 13182

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<400> 1655
cgtaccgcgc gtgtacgagg ctcttgaggt acaaaatggt agtgagaatc tgggtgctgga 60
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tctgcgtcgc ggtctggaag tcaaagacct cgagcatccg atcgaagtc cggtaggttaa 180
agcaacgctg ggctcgtatca tgaacgtact gggccaaccg gtagacatga aaggcgacat 240
cggcgaagaa gagcgttggg cgattcaccg cgcagcgcct tcctacgaag agttgtcaaa 300
ctctcaggaa ctgctggaaa ccggcatcaa agttatcgac ctgatgtgtc cgtttgcgaa 360
gggcggtaaa gttggtctgt tcggtggtgc ggggtgtaggt aaaaccgtaa acatgatgga 420
gctgatccgt aacatcgcca tcgagcactc cggttactcc gtgtttgcgg gcgtagggtga 480
acgtactcgt gagggtaacg acttctacca cgaaatgacc gactccaacg ttatcgataa 540
agtatccctg gtgtatggcc agatgaacga gccgcgggga aaccgtctgc gcgttgcgct 600
gaccggcctg accatggctg agaagttccg tgacgaaggt cgtgacgttc tgctgttcgt 660
cgataacatc tatcgttaca ccctggccgg tactgaagta tccgcactgc tgggtcgtat 720
gccttcagcg gtaggttacc agccgactct ggcggaagag atgggcgttc tgcagg 776
```

<210> 1656
 <211> 572
 <212> DNA
 <213> *Microsporum audouinii* ATCC 11347

```
<400> 1656
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aaaagtctcg aattagaaaa ttcttttcag atactaattt actatagaac aacattgcca 120
aggctcacgg tggttactcc gtcttcaccg gtgtcggaga gcgaaccctg gaaggaaacg 180
atctgtacca tgaaatgcag gaaactcgtg tcatccaact tgatggcgag tccaaggctc 240
ccctggtctt cggtcagatg aacgagcccc cagggtgccg tgcccgtgtt gctcttactg 300
gtttgaccat tgctgagtac ttccgtgatg aggaagggtca agacgggtatg ttcttttaaat 360
tagatatctt ctggagaaac agcgtctaac aaattcttcc agtgcttctc ttcatcgaca 420
acatcttccg tttcactcag gctgggttccg aagtgtctgc cctgcttggt cgtattccat 480
ctgccgtcgg ttaccaaccc actcttgccg tcgacatggg tggatatgcag gaacgtatta 540
ccaccaccaa gaagggtacc attacctccg tc                                     572
```

<210> 1657
 <211> 790
 <212> DNA
 <213> *Obesumbacterium proteus* ATCC 12841

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<400> 1657
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tcagcagcag ctaggcggcg gcgttgtagc ctgtatcgct atgggtactt ctgacgggtc 120
gcgtcgcgga ctggacgttg ttgacctgga gcacccgatt gaagtcccag taggtaaagc 180
gaccttaggc cgcattatga acgtactggg tgagccaatt gatatgaagg gtgatatcgg 240
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```

cgaagaagat cgctgggcta ttcaccgtga agctccaagc tacgaagaac tgtctaactc 300
gcaagaactg ctggaaaccg gtatcaaggt aatggacttg atttgtccgt tcgctaaggg 360
cggtaaaagtc ggtctgttcg gtgggtgcggg tgttggtaaa acagtaaaca tgatggagct 420
gatccgtaac atcgcgatcg agcactcagg ttactctgta tttgccggcg tgggtgaacg 480
tactcgtgag ggtaacgact tctaccacga aatgaccgac tccaacgta tggacaaagt 540
atcactgggt tatggccaga tgaacgagcc accaggaaac cgtctgcgcg ttgcgctgac 600
cggctctgact atggctgaga agttccgtga cgaaggctcg gacgtactgc tgttcacga 660
taacatctac cgttatacct tggccggtag cgaagtatct gactgctgg gtcgtatgcc 720
ttctgcggta gggtatcagc caacgctggc ggaagagatg ggtgttctgc aagaacgtat 780
cacctctacc                                     790

```

<210> 1658

<211> 622

<212> DNA

<213> *Paracoccidioides brasiliensis* ATCC 200443

<400> 1658

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tgtcttcatt caggagctta tcaacaacat cgccaaggcc cacggtggtt actccgtctt 60
cactgggtgtg ggagagcgca ctcgtagagg aaacgacttg tatcacgaga tgcaagagac 120
ttccggttacc cagctcgaag gcgaatccaa ggttgccctc gtcttcggtc aaatgaacga 180
gcctccgggt gctcgtgctc gtgttgctct caccggtcgt aagtgtcctt tcccagattt 240
ctcttcccca gtttctggac ccacttttcc cttccaccac cattctactg ggtaggacca 300
agatagcact gcctattctg gtgccttcc accgectact ctactgccta ttccaccacc 360
ttttctaccg cctcttctac ttgctattgt atactaactt actcaaacag ttactattgc 420
tgagtacttc cgtgacgctg agggccagga tgtgcttctc ttcacgcaga acattttccg 480
tttcacccag gccggttccg aggtgtccgc tcttctcggg cgtatccctt ccgccgtcgg 540
ttaccagccc acccttgccc tcgacatggg tggtatgcag gagcgtatca ccaccaccaa 600
gaagggatcc attacctccg tc                                     622

```

<210> 1659

<211> 794

<212> DNA

<213> *Plesiomonas shigelloides* ATCC 14029

<400> 1659

```

gacgctgtac ctcaggtgta cgatgcactg acagttgagg gtgctgagct ggtactggaa 60
gtgcagcagc agctgggtgg tgggtgttgt cgctgtatcg cgtatgggtg ctctgatggc 120
ctcaagcgcg gtctgaaagc gcacaatact ggtgctccta tctactgtacc ggtgggtgtg 180
gaaacactgg gccggatcat ggatgtgttg ggtaaccgca ttgaccagaa aggtccaatc 240
ggtgaacaag atcgctgggt gatccaccgt gaagcaccaa gctacgaaga tcaggctaac 300
agcactgaac tgctggaaac cgggtatcaag gttatcgacc tgggtatgcc gtttgcgaaa 360
ggcggtaaaag tcgggtctgt cgggtggtgcc ggtgtaggta aaaccgtaaa catgatggag 420
ctgatccgta acatcgcgat cgagcactcc ggttattccg tgtttgcggg cgtgggtgag 480
cgtaccctgt aaggtaacga cttctaccac gaaatgacag actccaacgt actggacaaa 540
gtatccctgg tgtacggtca gatgaacgag ccgccaggta accgtctgcg cgtagcactg 600
accggcctga ccattgcgga gaaattccgt gatgaaggct gtgacgtact gctgttcatc 660
gataacatct accgttatat cctggcgggg accgaagtat cggcactgct gggccgtatg 720
ccttctgcgg taggttatca gccaacgctg gcggaagaga tgggtgtact gcaagagcgt 780
attacctcta cccg                                     794

```

<210> 1660

<211> 799

<212> DNA

<213> *Shewanella putrefaciens* ATCC 8071

<400> 1660

```

aggtatatga cgctstgaag atcacaggtg aaggcgccctg taatggtttg gtgctggaag 60
ttcagcaaca gctaggcggg ggtgtagtct gtactatcgc tatgggttct tctgatggtc 120
tgctcgtggg tcttgagggt gttaactcag gttcacctat ttctgttctt gttggtagcc 180
ccacgcttgg ccgtatcatg aacgtattag gtgagcctat tgatgaagcg ggtccaatcg 240
gtgaagaaga gcgttatggt attcaccgtg cagcaccttc atatgaagat caatcgaaca 300
ctactgaact gttagagaca ggtatcaagg ttattgacct tgtttgtcca ttcgctaagg 360
gtggtaaagt aggtctgttc ggtgggtgcgg gtgttggtaa aacagttaac atgatggaac 420

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tgattaacaa	catcgctaaa	gctcactcgg	gtcttttcggt	gttcgcgcgg	gtgggtgaac	480
gtactcgtga	aggtaacgac	ttctactacg	agatgaaaga	ttctggcggt	ctcgacaaaag	540
tggccatggt	ttatggtcag	atgaacgagc	caccaggaaa	ccgtttacgc	gtagcactgt	600
cagggtctgac	aatggctgag	aagttccgtg	acgaaggctc	tgacgtattg	ttgttcggtg	660
acaacatcta	ccgttatacc	ttagccggta	ctgaagtatc	tgacgtgta	ggccgtatgc	720
cttctgcggg	aggttatcaa	ccaacattgg	ctgaagaaat	gggcggttctg	caagagcgta	780
ttacttcaac	taagacggg					799

<210> 1661

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1661

tggaagcgga aaatcctg

18

<210> 1662

<211> 774

<212> DNA

<213> Campylobacter curvus ATCC 35224

<400> 1662

ctatgcctca	aactagagag	catatcttgc	tatctcgcca	agtaggcgtt	ccatatatcg	60
ttgtatttat	gaacaaagcc	gatatggctg	atgacgctga	gcttcttgag	ctagtcgaga	120
tggaaattcg	cgagcttctt	aacgagtaca	acttccctgg	cgatgatact	cctatcatat	180
caggttctgc	tcttaaagcc	ctcgaagagg	ctaaagcagg	cggtgatggc	gagtggtcag	240
caaaagttct	tgagcttatg	gataaagtcg	atgagtatat	cccaactcca	gttcgtgcta	300
ccgataaaga	cttcctgatg	cctatcgaag	acgtttttct	tatctcaggt	cgtggaacgg	360
tcgttactgg	taggatcgaa	aaagggtgct	taaaagttgg	cgatactatc	gagatcgttg	420
gtatcaaacc	tactcaaact	acgacagtta	ctggcggtga	gatgtttagg	aaagagatgg	480
aacaaggcga	ggccgggtgat	aacgtaggtg	ttcttttaag	aggtactaaa	aaagaagacg	540
tcgagcgcg	catggttctt	tgtaaagccaa	aatcaatcac	tcctcataca	aaatttgagg	600
gtgaggttta	tatcctaaca	aaagaggaag	gcggacgcca	cactccattc	tttaacaact	660
atagaccaca	attttatgta	agaacaacag	acgttacagg	ttctatcaca	cttcagaag	720
gaactgagat	ggttatgcct	ggagataatg	tcagaatttc	cggtgaactc	atcg	774

<210> 1663

<211> 791

<212> DNA

<213> Campylobacter rectus ATCC 33238

<400> 1663

ttctgcggct	gacggcccaa	tgccacaaac	tagagagcac	atcttgctat	ctcgccaagt	60
aggcggttcc	tatatcggtg	tttttatgaa	caaagccgat	atggtcgatg	atgccgagct	120
tcttgagctg	ggtgagatgg	agattcgcga	gcttctaaac	gagtatgatt	tccctgggtga	180
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agaaggcgag	tggtctgcaa	aaattcttga	gcttatggct	aaagttgacg	agtatatccc	300
gactccgggt	cgtgcaacgg	ataaagactt	cttgatgcct	attgaggacg	ttttctctat	360
ctccgggtcgc	ggcaccgtcg	ttaccggcag	aatcgaaaaa	ggtatcgtaa	aagtcgggtga	420
tactatcgag	atcgtaggta	tccgcgatac	tcaaacaaact	accgttaccg	gcgttgagat	480
gttcagaaaa	gagatggatc	aaggcgaagc	gggcgataac	gtaggcggtc	ttctaagagg	540
cactaaaaaa	gaagacggtg	agcgcggtat	ggttctttgc	aaacctaata	caatcactcc	600
tcacactaaa	tttgagggag	aggttttatat	cttaactaaa	gaggaaggcg	gacgccatac	660
tccattcttt	aataactata	gaccgcagtt	ttatgtaaga	actaccgacg	ttaccgggtc	720
tatcactctt	ccggaaggaa	cagagatggt	tatgcctggc	gataacttaa	agataagcgt	780
tgagcttatac	g					791

<210> 1664

<211> 810
<212> DNA
<213> *Fonsecaea pedrosoi* ATCC 18831

<400> 1664
cgacggacag atgccccaga ccaggaggca cttgctcctc gcccgccagg tcggtgtcaa 60
gcgcattgtc gtcttcgtca acaagggtcg tgccattgag gacaaggaga tgttggagct 120
cgtcgagatg gagatgcgtg agcttctctc cagctacggc ttcgaggggtg acgacactcc 180
catcgtcatg ggttccgccc tttgcgccat tgagggccgc gagcccgaca ttggtgtcga 240
gaagattgac gagctcctcg agcacgtcga cacctggatc cccacccccg agcgtgacat 300
cgccaagcct ttcctcatgt ccgttgagga cgtcttctcc attcccggcc gtggtaccgt 360
cgcttctggc cgtgtcgagc gtggtgtcct gaagaaggat tccgaagtcg agcttgtcgg 420
caagaacaag aacccccatca agaccaagggt taccgacatc gagaccttca agaagtcttg 480
cgacgagtcc cgcgctgggtg acaactccgg tctccttctc cgtggtgtca agcgtgacga 540
tgtcctccgt ggcattgggtc ttgtccagcc cggcaccacc aaggcccaca agaagttcct 600
tgcctccatg tacgtcctca ccaaggagga ggttgccgc cactactggt tcgccaacaa 660
ctacaagccc cagatgttca tccgtaccgc cgatgaggcc gccactctta cctggcccga 720
gggtaccgag gaggacaaga tggtcattgcc cggtgacaat gtcgagatga tctgcgagat 780
ccacaagccc attgccgtcg agcaaggcca 810

<210> 1665
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1665
cagtacaggt agacttctg 19

<210> 1666
<211> 888
<212> DNA
<213> *Microsporum audouinii* ATCC 11347

<400> 1666
atgattgcga aacctacttg ctgtggaaga atttggatat tctaacattt ctctaggcct 60
caaaccagag agcatctgct ccttgcccgc cagggtcggtg ttcagaagct cgtcgttttc 120
gttaacaagg tcgacgctgt tgaggacca gagatgttgg aacttgtcga gctagagatg 180
cgtgagctgc tcagccacta tggtttcgag ggtgaggaga cccaatcat ttttggctct 240
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ttgaacgctg tggatacctg gattcccacc ccagagcgtg cactgataa gcctttcctt 360
atgtccattg aggaagtttt ctccatctct ggctcgtggtt ccgtcgtttc cggctcgtgtc 420
gagcgtggta tcctcaagaa ggactctgat gtcgaaattg tgggtggatc tgatacaccc 480
atcaagacga aggtcaccga cattgaaacc ttcaagaagt cttgtgacga atcccagct 540
ggtgacaact ccggtctact tctccgaggt gtcaagcgtg aggacttgag acgtggaatg 600
gttgttgctg ctcccggatc gaccaaggct cataccgact tcatggtctc cctttatgtt 660
ctgaccgagg ctgagggtgg ccgttccaat ggattcacc acaagtaccg cccacagatg 720
ttcatccgta ctgccggtat gtaaaccctt tttctacat tcaactttgt tcaccactga 780
cttgataact ttaccgcaga cgaagccgca tctttcagct ggcctggaga ggatcaagac 840
aagaaggcca tgcctggtga caatgtcgag atgatttgca agaccctc 888

<210> 1667
<211> 793
<212> DNA
<213> *Piedraia hortai* ATCC 24292

<400> 1667
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gttttcgtca acaaggttga tgctatcgac gacccggaga tgctggagct tgtcgagatg 120
gagatgcgtg aacttctcag cacatacgggt ttcgaggggt acgagacccc tgttattatg 180

ggctccgcgc	tcatggctct	caacaaccag	cgccccgaga	ttgggtcaaca	gaagattgat	240
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ttcctgatgt	ctgttgagga	tgtcttctcc	attgctggcc	gtgggtaccgt	tgtgtccggc	360
cgtgtggagc	gcggtaccct	caagcgtgat	gaggaagtcg	agcttgtcgg	caagggtgtc	420
gaccccatca	agaccaaggt	caccgatatc	gagactttca	agaagtcctg	cgaggaggct	480
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tacgtttctca	ccaaggagga	gggtgggtcgc	cacactgggt	tcggcgagca	ctaccgtccc	660
cagctctacc	tccgtacctc	agacgagtct	gtcgatctga	ccttccccga	gggaactgag	720
gatcaccact	ccaagatcgt	catgcctggg	gacaacatcg	agatggtcgt	cacgatgact	780
cacgccaaacg	cta					793

<210> 1668
 <211> 891
 <212> DNA
 <213> Escherichia coli strain K-12 KL1699

<400> 1668						
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ggcccgatgc	cgcgactcgc	tgagcacatc	ctgctgggtc	gtcaggtagg	cgttccgtac	120
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gttcgtgggt	ctgctctgaa	agcgtgggaa	ggcgacgcag	agtgggaagc	gaaaatcctg	300
gaactggctg	gcttccctgga	ttcttacatt	ccggaaccag	agcgtgcgat	tgacaagccg	360
ttcctgctgc	cgatcgaaga	cgtattctcc	atctccgggtc	gtgggtaccgt	tggtaccggt	420
cgtgtagaac	gcggtatcat	caaagtgtgt	gaagaagtgt	aaatcgttgg	tatcaaagag	480
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gctgggtgaga	acgtagggtg	tctgctgcgt	ggtatcaaac	gtgaagaaat	cgaacgtggg	600
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<210> 1669
 <211> 805
 <212> DNA
 <213> Saksenaea vasiformis ATCC 60625

<400> 1669						
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tcccgggcga	cgacgttccc	gtcgtcaagg	tctccgtctc	gaaggccctc	gagggcgaca	240
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ccgagcgtga	cgtcgacaag	ccgttcctca	tgccgatcga	ggacgtcttc	acgatcaccg	360
gtcgcggtac	ggtcgtcacc	ggccgtatcg	agcgtgggtg	cctgaagggtc	aacgagaccg	420
tcgacatcat	cggcatacaag	accgagaaga	ccaccaccac	ggtcaccggc	atcgagatgt	480
tccggaagct	cctcgacgag	ggccaggccg	gtgagaacgt	cggctctgctc	ctccgtggca	540
tcaagcgcca	ggacgtcgag	cgccggccagg	tcatcatcaa	gcccgggctcg	gtcacgccgc	600
acacggagtt	cgaggcgcag	gcctacatcc	tgtccaagga	cgagggtggc	cgccacacgc	660
cgttcttcaa	caactaccgc	ccgcagttct	acttccgtac	gacggacgtg	accggcgtgg	720
tgaccctccc	cgagggcacc	gagatgggtca	tgccgggtga	caacaccgag	atgaagggtg	780
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<210> 1670
 <211> 935
 <212> DNA
 <213> Trichophyton tonsurans ATCC 56185

<400> 1670						
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ctaactatcag	tctaggcctc	agaccagaga	acatttgctc	cttgcccgcc	aggtcgggtg	120

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ccagaagctg gtcggttttcg ttaacaaggt cgatgccgtt gaggaccag agatgttgga 180
gcttgtcgaa cttgaaatgc gtgaactcct cagccactac ggtttcgagg gtgaggagac 240
ccccatcatt tttggctctg ctctctgtgc cctcgagtcg cgtecgacctg agcttgggtgt 300
cgagaagatt gacgagctat tgaacgccgt cgacacctgg atccccaccc cagagcgcg 360
cactgataag cctttcctca tgtccattga ggaagtgttc tctatctctg gtcgtggtac 420
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tgggtggtct accaccccta tcaagaccaa ggtcacccgat atcgaaacct tcaagaagtc 540
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caagtaccgc ccccaaattg tcatccgtat tgctggtag taaccaagt tcccgctatt 780
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aagaccctcc accccattgc tgccgaggct ggcca 935

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<210> 1671
 <211> 772
 <212> DNA
 <213> *Enterobacter aerogenes* ATCC 13048

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<400> 1671
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tggaagttaa agaccttgag cacccgatcg aagtccccgt aggtaaagcg actctgggcc 180
gtatcatgaa cgtcctgggt cagccgatcg acatgaaagg cgacatcggc gaagaagaac 240
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tggaaccggg catcaaagtt atcgacttga tgtgtccgtt cgctaaggggc ggtaaagttg 360
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tcgcgatcga gcaactccgt tactccgtgt ttgcgggctg tggtagcgt actcgtgagg 480
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gttacacctt ggccggtact gaagtatctg cactgctggg ccgtatgcct tcagcggtag 720
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<210> 1672
 <211> 1401
 <212> DNA
 <213> *Bordetella pertussis* strain Tohama 1

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<400> 1672
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aaggccaaga aactccaata a

1401

<210> 1673

<211> 797

<212> DNA

<213> Arcanobacterium haemolyticum ATCC 9345

<400> 1673

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accgtccaca	gttctacttc	cgtaccacgg	atgttacccg	cgtgatcacc	cttccagagg	720
gcaccgaaat	ggttatgcca	ggcgacaaca	ccgacatgac	agttgagctc	atccagccaa	780
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<210> 1674

<211> 785

<212> DNA

<213> Butyrivibrio fibrisolvens ATCC 19171

<400> 1674

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gtccaaccac	caagaccact	gtaactggcg	ttgaaatggt	ccgtaagtta	ctagacgaag	480
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tttacgtact	aagcaaggat	gaagggtggc	gtcacactcc	attcttcaag	ggctaccgtc	660
cacagttctt	cttccgtaca	accgatatta	ccggttctat	cgatctgaaa	gagggcgtag	720
agatggtaat	gccagggtgat	aacaccgaca	tgaccgtaac	cctaattccac	ccagtagcta	780
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<210> 1675

<211> 835

<212> DNA

<213> Campylobacter jejuni subsp. doylei ATCC 49349

<400> 1675

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tgaagggtgg	agacatactc	cattctttta	caactataga	ccacagtttt	atgtaagaac	720
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aaatgtgaga attactgtaa gcttgatcgc tccagtagca cttgaagaag gaact 835

<210> 1676

<211> 812

<212> DNA

<213> *Campylobacter lari* ATCC 43675

<400> 1676

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aactccggct	cgtgatacag	ataaagattt	cttgatgcca	atcgaagatg	ttttctcaat	360
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tattaaactt	gcagaaggcg	ttgagatggg	tatgccaggc	gataatgata	gaattactgt	780
aagtcttatt	gctccagttg	cacttgagga	ag			812

<210> 1677

<211> 828

<212> DNA

<213> *Campylobacter sputorum subsp. sputorum* ATCC 35980

<400> 1677

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gatgatgctg	agcttataga	gttgggtgaa	gttgagggtta	gagatttatt	aaatgaatat	180
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actggtgttg	aaatgtttag	aaaagagcta	gatcaagggtg	aagctggaga	taatgttggt	540
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gatgtaacag	gttctatata	tcttcctgag	ggaacagaga	tggatatgcc	tggtgataat	780
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<210> 1678

<211> 821

<212> DNA

<213> *Campylobacter upsaliensis* ATCC 49815

<400> 1678

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atgttttttc	aattttctggt	cgtggaactg	ttgtaacagg	tagaattgaa	aaagggtgtg	420
ttaaagtcgg	cgatactatt	gagatagtag	gtatcaaaga	tactcaaact	acaacagtta	480
caggcggttg	gatgttttaga	aaagaaatgg	atcaagggtga	ggctggcgat	aatgtcgggtg	540
tgcttttaag	aggaacaaaa	aaagaagatg	ttcttcgtgg	tatggttctt	gcaaagccta	600
gaattctatca	ctcctcatac	tgatttttgaa	gcagaagttt	atattctaaa	taaagatgag	660
ggcggtcgcc	atactccttt	ctttaacaat	tatcgtccgc	agttttatgt	aagaacgact	720
gatgtaactg	gttctattaa	attagctgat	gggtgtgaga	tggttatgcc	gggtgaaaaat	780

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821

<210> 1679

<211> 783

<212> DNA

<213> Globicatella sanguis ATCC 51173

<400> 1679

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caa						783

<210> 1680

<211> 823

<212> DNA

<213> Lactobacillus acidophilus ATCC 4356

<400> 1680

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<210> 1681

<211> 793

<212> DNA

<213> Leuconostoc mesenteroides subsp. dextranicum ATCC 19255

<400> 1681

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793

<210> 1682
<211> 796
<212> DNA
<213> *Prevotella buccalis* ATCC 35310

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<210> 1683
<211> 800
<212> DNA
<213> *Ruminococcus bromii* ATCC 27255

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<210> 1684
<211> 545
<212> DNA
<213> *Paracoccidioides brasiliensis* ATCC 32075

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caagccggtt ccgaagtgtc cgccctgcta ggccgcatcc cctccgccgt cggctatcaa 480
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<210> 1685

<211> 1020
<212> DNA
<213> *Candida norvegica* ATCC 36586

<400> 1685
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gaaccacctt caagaccaac tgaaaaacca ttgagattgc cattgcaaga tgtctacaag 540
attgggtggta tcggaaccgt accagtcggt actactgaag tcaaactctgt tgaaatgcat 600
atgattgtta ctttcgcccc agccggtggt actactgaag tcaaactctgt tgaaatgcat 660
cacgaacaat tagaagctgg ttaccaggt gacaatgttg gtttcaacgt caagaatggt 720
tcagttaaag aaatcagaag aggttaahgtt gctggtgact ccaagaacga tccacaaaaa 780
ggtgctgaat ctttcaacgc tcaagttatt gtcttgaacc atccagggtca aatctytgct 840
ggttactytc cagtttttggg ttgtcacact gccacattg cttgtaaatt cgatgaaatc 900
ttggaaaaga ttgacagaag atccggtaag aaattggaag aaaatccaaa attcatcaaa 960
tctggtgacg ctgctawtgt caaatttgtt ccatwtaaac cattrtgtgt tgaagctttc 1020

<210> 1686
<211> 929
<212> DNA
<213> *Aspergillus nidulans* strain WSA-176

<400> 1686
tgctgcttcc gatggtcaaa tgtacgattg atattccttc cagccagtca ggataacagc 60
tgataccagt tgcaaatagg ccccagactc gtgagcactt gttgcttgcc cgtcagggtg 120
gtgtccagaa gatcgttgct ttcgtcaaca aggttgacgc tgtcgatgac cctgagatgt 180
tgagagcttg tgagctcgag atgcgtgagc tctcaacac ttacggtttc gagggagagg 240
agacccctat catcttcggt tccgccctgt gcgctctcga aggccgccgc gaggacattg 300
gtactcagcg tattgactcc ctctcagagg ccgttgacac ttggatccct accccccagc 360
gtgacttggg caagcccttc ctgatgtcca ttgaggaagt tttctccatt ggtggctcgtg 420
gtaccgtcgc ctctggctgt gtcgagcgtg gtctcctcaa gaaggatacc gaagttgaaa 480
ttcacggtgc tgatggtatt ctgaagacca aggtcaccca cattgagacc ttcaagaaga 540
gctgcgatga gtctcgtgct ggtgacaact ccggtcttct cctccgtggt atccgtcgtg 600
aggatgttcg tcgtgggtatg gtcacgcgtg cccctggctc catcaaggcc tccaagaagt 660
tcatggtctc catgtacgtc ttgactgagg ctgaagggtg ccgcaagaac ggcttcggtg 720
ccaactaccg cccccaggct ttcacccgca ctgctggtaa gtttcgaact atttgattca 780
ttgatcacgt ccctaactgt tacttttagac gaggcttgcg accttcattt ccctgatgag 840
gccgacaagg accgccagct catgcccggg gacaacgtcg aaatggctct caacctcaac 900
aaccctggtg ctgctgaggc tggacagcg 929

<210> 1687
<211> 951
<212> DNA
<213> *Aspergillus terreus* strain WSA-174

<400> 1687
tgccgcttcc gatggtcaga tgtacgctca agccccagtt tccatataaa cataaacgat 60
ctatcatcag cacaacgctg acttcttcgc ttccaggccc cagaccctg agcacttgct 120
gttggcccggt caggctcggtg tccagaagat cgtggctctc gtcaacaagg tcgatgccgt 180
tgatgaccgg gagatgttgg agctcgttga gctggaaatg cgcgagctcc tgaccagcta 240
cggattcgag ggtgaagaga cccccatcat ctccggttct gctctctgag ctcttgaggg 300
ccgccgtcct gagattggta ctgagaagat tgacgagctg atgcacgccg tcgacacctg 360
gatccccacc cccagcgtg acctcgacaa gcccttctct atgtccgtcg aggaagtctt 420
ctccatttgt ggtcgtggta ccgtcgtctc cggccgtgct gagcgtggta ttctgaagaa 480
ggatagcgaa gtcagatca tcggtggtgc ttctgacgcc acgaagacca aggtcactga 540
catcgagacc ttcaagaagt cttgcgacga gtctcgcgct ggtgacaact ctggtctcct 600
cctccgtggt atccgtcgtg aggatgttcg gcgtgggtat gtcattgctg ctctggcgag 660
caccaaggcc caccagaagt tcttctctc tatgtacgtc ctactgagg ctgagggtg 720

-568-

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ccgtcgtacc ggattcggta ccaactaccg cccccaggtc ttcattccgta ctgccggtaa 780
gtgttcctgg aagaggcttt gagcctatat aggatctcgg ataatttact aatccaccat 840
atagatgagg ccgctgacct cagcttcccc gacaacgatg actcccgcg tgatcatgcc 900
ggtgacaacg ttgagatggt cctgaagacc caccgccccg tggctgctga g 951
```

<210> 1688
<211> 823
<212> DNA
<213> *Candida norvegica* ATCC 36586

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<400> 1688
cggtgccgct accgatggtc aaatgcctca aactagagaa catttgctat tggctagaca 60
ggttggtggt caacacattg tcgtgtttgt taacaagggt gacactattg atgatccaga 120
aatggttgaa ttggttgaaa tggaaatgag agagttgatt gccacttatg gtttcgatgg 180
tgataacacc ccagttatca tgggttctgc tctatgtgct ttggaagggtc gtgaacctga 240
aatcgggtgct caatcaatcg acagattggt ggaagccgtt gatgaatata ttccaactcc 300
aactagagat ttggaaaaac cattcttgat ggggtgttgaa gatgtcttct ccatttctgg 360
tagagggtacc gtctgtaccg gtcgtgttga aagaggtaac ttgaagaaag gtgatgaaat 420
cgaaattgtc ggctacaaca agactccaat caaaaccacc gtcaccggtt ttgagatggt 480
caaaaaggaa ttgaccaaga ctatggctgg tgataactgt ggtatcttat tacgtggtgt 540
taagagagat gatatacaaga gaggtatggt tatctctaaa gtcaacaccg tttccgcaca 600
caccaaattc ttggcctctt tatacgtctt gactaaagaa gaagggtggc gtcattcagg 660
ttttgctgaa aactacagac ctcaattggt catcagaacc ggtgatgtca ctgttacttt 720
aaccttcccc gaagatgctg atcactctca gcaagtctta ccagggtgaca acgttgaaat 780
ggaatgtacc ttgggttcac caactgctct tgaaaccggt caa 823
```

<210> 1689
<211> 803
<212> DNA
<213> *Candida parapsilosis* ATCC 201076

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<400> 1689
gctgctactg acggtcaaat gcctcaaact agggaaacata tgttggttggc gagacaagtt 60
ggtatccaaa acttggttgt ttttggttaac aaagttgata ccattgatga cccagaaatg 120
ttggaattgg ttgaaatgga aatgagggaa ttattgagct cttatgggtt tgatgggtgaa 180
aacactccag ttatcatggg atcagccttg tgtgctttag aaggtaaaca accagaaatc 240
ggtgttcaag ccattcaaaa attattggat gctgttgatg aatatattcc aactccagaa 300
agagatgctg accaaccatt tttgatgcca gtggaagatg tgttttctat ttcaggtaga 360
ggaaccggtt tcaccggaag agttgaaaga ggtatgttga agaaagggtg agaagtakaa 420
gtcattgggtg aaaactcatt taaggctact tccacgggta ttgagatggt caaaaaggaa 480
ttggatgccg ctatggcccg tgacaactgt ggtattttgt tgagagggtg caagagagac 540
gaagtcaaga ggggtatggt tttggccaaa ccaggtaacca ccacccaca ccaaaagttt 600
ttggcttcca tttatatctt gactgctgaa gaagggtggac gtagtacccc tttcagttaa 660
ggatacaaac cacaatgttt ctttagaact agtgatgtta ccacgacatt tactttccca 720
gaagggtgaag gtgttgacca ctcacaaatg gttatgccag gagrcaatgt tgaaatggtg 780
ggaactttta tcaagaaagc tcc 803
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<210> 1690
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1690
caggctcgtg tgcgactgaa gaa

23

<210> 1691
<211> 25
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1691

cacagataaa cctgagtgtg ctttc

25

<210> 1692

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1692

ggtgagaact gtggatatctt actt

24

<210> 1693

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1693

catttcaacg ccttctttca actg

24

<210> 1694

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1694

aaggcaagga tgacaacggc

20

<210> 1695

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1695

acgatttcca cttcttctg g

21

<210> 1696

<211> 20

<212> DNA

<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1696
atgttcctgt agttgctgga 20

<210> 1697
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1697
tttcttcagc aataccaaca ac 22

<210> 1698
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1698
ggaatcaaca gatgggtttac aaa 23

<210> 1699
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1699
gcatcttctg ggaaaggtgt 20

<210> 1700
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1700
aagatgcgga aagaagcgaa 20

<210> 1701
<211> 23
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1701
attatggatc agttcttggg tca

23

<210> 1702
<211> 213
<212> DNA
<213> Streptococcus gordonii strain Challis V288

<400> 1702
ttcatagacg ctgagcacgc tttggatcca tcttacgcgg ctgctctagg tgtaaatatt 60
gatgagctgt tgctatctca accagattct ggtgagcaag gtttagaaat tgcaggaaaa 120
ttgattgact ctggggcagt tgatttagtt gtcacgact ctggtgcagc tcttgtacca 180
cgtgcggaaa tcgatggaga tatcgggtgat agc 213

<210> 1703
<211> 692
<212> DNA
<213> Streptococcus mutans strain GS-5

<400> 1703
gggcccgaat cttctggtaa gacaactgtc gctcttcatg ctgctgctca ggcgcaaaaa 60
gatggcggta ttgccgcttt cattgatgca gaacatgcc ttgatccagc ctatgctgct 120
gctcttggcg ttaatatattga tgagcttttg ctttcacaa cagattcagg agaacagggt 180
cttgaaattg cagggaaatt gattgattct ggcgctgttg atttagttgt tgttgactca 240
gtggcagctt tagtaccacg tgcggagatt gacggagata ttggtaatag tcatgttggc 300
ttacaagcac gcatgatgag tcaagcgatg cgtaaattat cagcttcaat caataaaaca 360
aaaaccattg ctatttttat taatcaattg cgggaaaaag ttggtattat gtttggtaat 420
ccagaaacaa cccctggcgg gcgtgccttg aagttttatt cttctgtgcg tcttgatgtc 480
cgcggaata ctcaaattaa aggaaccggg gaacaaaaag acagcaatat tggtaaagag 540
accaaaatta aagttgttaa aaataaagtt gctccaccat ttaaggaagc tttttagtaa 600
attatatatg gtgaaggcat ttctcgtaca ggtgaattag ttaagattgc cagtgatttg 660
ggaattatcc aaaaagctgg agcttggtac tc 692

<210> 1704
<211> 1204
<212> DNA
<213> Streptococcus pneumoniae

<400> 1704
atggcgaaaa aaccaaaaaa attagaagaa atttcaaaaa aatttggggc agaacgtgaa 60
aaggccttga atgacgctct taaattgatt gagaaagact ttggtaaagg atcaatcatg 120
cgtttgggtg aacgtgcgga gcaaaagggt caagtgatga gctcagggtt ttagctctt 180
gacattgccc ttggctcagg tggttatcct aaggggacgt tcatcgaaat ctatggccca 240
gagtcactct gtaagacaac ggttgccctt catgcagttg cacaagcgca aaaagaagg 300
gggattgctg cttttatcga tgcggaacat gcccttgatc cagcttatgc tgcggccctt 360
ggtgtcaata ttgacgaatt gctcttgtct caaccagact caggagagca aggtcttgag 420
attgcgggaa aattgattga ctcagggtga gttgatcttg tcgtagtcga ctcagttgct 480
gcccttgctt ctcgtgcgga aattgatgga gatatcggag atagccatgt tggtttgag 540
gctcgtatga tgagccaggc catgcgtaaa cttggcgctt ctatcaataa aacaaaaaca 600
attgccattt ttatcaacca attgcgtgaa aaagttaggag tgatgtttgg aaatccagaa 660
acaacaccgg gcggacgtgc tttgaaattc tatgcttcag tccgcttgga tgttcgtggt 720
aatacacaaa ttaagggaac tggatgatca aaagaaacca atgtcggtaa agaaactaag 780
attaaggttg taaaaaataa ggtagctcca ccgtttaagg aagccgtagt tgaaattatg 840
tacggagaag gaattttctaa gactggtgag cttttgaaga ttgcaagcga tttggatatt 900
atcaaaaaag caggggcttg gtattcttac aaagatgaaa aaattgggca aggttctgag 960
aatgctaaga aataacttggc agagcaccga gaaatctttg atgaaattga taagcaagtc 1020
cgttctaaat ttggcttgat tgatggagaa gaagtttcag aacaagatac tgaaaaacaa 1080
aaagatgagc caaagaaaga agaagcagtg aatgaagaag ttccgcttga cttaggcgat 1140
gaacttgaag tcgaaattga agaataagct gttaaagcag tggagaaatc cgctactttt 1200
tcga 1204

<210> 1705
 <211> 981
 <212> DNA
 <213> *Streptococcus pyogenes* strain NZ131

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<400> 1705
atgCGttcag gaagtctagc tcttgatatt gcttggatag ctggtgggta tcctaaagga 60
cgtatcatcg aaatctatgg tccagagtct tccggtaaaa cgactgtggc tttacatgct 120
gtagcacaag ctcaaaaaga aggtggaatc gcagccttta tcgatgccga gcatgcgctt 180
gatccagctt atgctgctgc gcttgggggtt aatattgatg aacttctctt gtctcaacca 240
gattctggag aacaaggact tgaaattgca ggtaaattga ttgattctgg tgcgggttgac 300
ctggttggtg tcgattcagt agcagcttta gtgccacgtg ctgaaattga tggatgatatt 360
ggcgaatagc atgtcggatt gcaagcacgt atgatgagtc aggccatgcg taaattatca 420
gcttctatta ataaaaacaa aactatcgca atctttatca accaattgcg tgaaaaagtt 480
ggtgtgatgt ttggaaatcc tgaaacaaca ccagggtggc gagctttgaa attctatgct 540
tctgttcggc tggatgtgcg tggaaacaac caaattaaag gaactgggta ccaaaagata 600
gccagcattg gtaaggagac caaatcaag gtgtgtaaaa acaaggctgc tccgccattt 660
aaggtagcag aagttgaaat catgtatggg gaaggtatct ctcgtacagg ggagcttggt 720
aaaattgctt ctgatttggc cattatccaa aaagcagggt cttgggttct ttataatggt 780
gagaagattg gccaaagggt tgaaaaatgct aagcgttatt tggccgatca tccacaattg 840
tttgatgaaa tcgaccgtaa agtacgtgtt aaatttggtt tgcttgaaga aagcgaagaa 900
gaatctgcta tggcagtagc atcagaagaa accgatgatc ttgctttaga tttagataat 960
ggtattgaaa ttgaagatta a                                     981
```

<210> 1706
 <211> 312
 <212> DNA
 <213> *Streptococcus salivarius* subsp. *thermophilus*

```
<400> 1706
gcgtatgcac gagctctagg tgtaatatc gatgagcttc ttttgtcgca gcctgattct 60
ggtgagcaag gtctcgaaat tgcaggtaag ctgattgact ctgggtgcagt ggatttagtt 120
gttgttgact cagttgcgcc cttcgtacca cgtgcagaaa ttgatggaga tagtgggtgac 180
agtcattgat gacttcaagc gcgtatgatg agtcaagcca tgcgtaaact ttctgcatct 240
attaataaaa caaaaacgat tgctatcttt attaaccagt tgcgtgaaaa agttgggtatc 300
atgtttggta ac                                     312
```

<210> 1707
 <211> 831
 <212> DNA
 <213> *Escherichia coli*

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<400> 1707
atgaaaaaca caatacatat caacttcgct attttttttaa taattgcaaa tattatctac 60
agcagcgcca gtgcatcaac agatatctct actgttgcat ctccattatt tgaaggaaact 120
gaaggttggt ttttacttta cgatgcatcc acaaacgctg aaattgctca attcaataaa 180
gcaaagtgtg caacgcaaat ggcaccagat tcaactttca agatcgcatc atcacttatg 240
gcatttgatg cggaataaat agatcagaaa accatattca aatgggataa aacccccaaa 300
ggaatggaga tctggaacag caatcataca ccaaagacgt ggatgcaatt ttctgttggt 360
tggttttcgc aagaaataac ccaaaaaatt agattaaata aaatcaagaa ttatctcaaa 420
gattttgatt atggaaatca agacttctct ggagataaag aaagaaacaa cggattaaca 480
gaagcatggc tcgaaagtag cttaaaaatt tcaccagaag aacaaattca attcctgcgt 540
aaaattatta atcacaatct cccagttaaa aactcagcca tagaaaacac catagagaac 600
atgtatctac aagatctgga taatagtaca aaactgtatg ggaaaactgg tgcaggattc 660
acagcaaata gaaccttaca aaacggatgg tttgaagggt ttattataag caaatcagga 720
cataaatatg tttttgtgtc cgcacttaca ggaaacttgg ggtcgaattt aacatcaagc 780
ataaaagcca agaaaaatgc gatcaccatt ctaaacacac taaatttata a                                     831
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<210> 1708
 <211> 846
 <212> DNA
 <213> *Enterococcus faecalis* strain HH22

<400> 1708
ttgaaaaagt taatatTTTT aattgtaatt gcttttagttt taagtgcattg taattcaaac 60
agttcacatg ccaaagagtt aaatgattta gaaaaaaaat ataatgctca tattgggtgtt 120
tatgcttttag atactaaaag tggtaaggaa gtaaaattta attcagataa gagattttgcc 180
tatgcttcaa cttcaaaaagc gataaatagt gctattttgt tagaacaagt accttataat 240
aagttaaaata aaaaagtaca tattaacaaa gatgatatag ttgcttattc tcctatttta 300
gaaaaaatatg taggaaaaga tatcacttta aaagcactta ttgagggttc aatgacatat 360
agtgataata cagcaaacaa taaaattata aaagaaatcg gtggaatcaa aaaagttaaa 420
caacgtctaa aagaactagg agataaagta acaaatccag ttagatatga gatagaatta 480
aattactatt caccaaagag caaaaaagat acttcaacac ctgctgcttt cggttaagact 540
ttaaataaac ttatcgcaaa tggaaaatta agcaaagaaa acaaaaaatt cttacttgat 600
ttaatgttaa ataataaaag cggagatact ttaattaaag acggtgttcc aaaagactat 660
aagggttgctg ataaaagtgg tcaagcaata acatatgctt ctagaaatga tgttgctttt 720
gtttatccta agggccaatc tgaacctatt gtttttagtca tttttacgaa taaagacaat 780
aaaagtgata agccaaatga taagttgata agtgaaaccg ccaagagtgt aatgaaggaa 840
ttttaa 846

<210> 1709
<211> 555
<212> DNA
<213> *Pseudomonas aeruginosa*

<400> 1709
atgtccgcga gcaccccccc cataactctt cgcctcatga cccgagcgcga cctgccgatg 60
ctccatgact ggctcaaccg gccgcacatc gttgagtggt ggggtggcga cgaagagcga 120
ccgactcttg atgaagtgtt ggaacactac ctgcccagag cgatggcgga agagtccgta 180
acaccgtaca tcgcaatgct gggcgaggaa ccgatcggct atgctcagtc gtacgtcgcg 240
ctcggaagcg gtgatggctg gtgggaagat gaaactgac caggagtgcg aggaatagac 300
cagtcctctg ctgacccgac acagttgaac aaaggcctag gaacaaggct tgtccgcgct 360
ctcgttgaac tactgttctc ggacccacc gtgacgaaga ttcagaccga cccgactccg 420
aacaaccatc gagccatacg ctgctatgag aaggcaggat tcgtgcggga gaagatcatc 480
accacgcctg acgggcccgc ggtttacatg gttcaaacac gacaagcctt cgagagaaaag 540
cgcggtgttg cctaa 555

<210> 1710
<211> 732
<212> DNA
<213> *Staphylococcus aureus*

<400> 1710
atgaaccaga aaaaccctaa agacacgcaa aatttttatta cttctaaaaa gcatgtaaaa 60
gaaatattga atcacacgaa tatcagtaaa caagacaacg taatagaaat cggatcagga 120
aaaggacatt ttaccaaaga gctagtcaaa atgagtcgat cagttactgc tatagaaatt 180
gatggaggct tatgtcaagt gactaaagaa gcggtaaacc cctctgagaa tataaaagtg 240
attcaaacgg atattctaaa attttccttc ccaaaacata taaactataa gatatatggt 300
aatattcctt ataacatcag tacggatatt gtcaaaaagaa ttacctttga aagtcagggt 360
aaatatagct atcttatcgt tgagaaggga tttgcgaaaa gattgcaaaa tctgcaacga 420
gctttgggtt tactattaat ggtggagatg gatataaaaa tgctcaaaaa agtaccacca 480
ctatattttc atcctaagcc aagtgtagac tctgtattga ttgttcttga acgacatcaa 540
ccattgattt caaagaagga ctacaaaaag tatcgatctt ttgtttataa gtgggttaaa 600
cgtgaatatc gtgttctttt cactaaaaac caattccgac aggcctttgaa gcatgcaaat 660
gtcactaata ttaataaact atcgaaggaa caatttcttt ctattttcaa tagttacaaa 720
ttgtttcact aa 732

<210> 1711
<211> 738
<212> DNA
<213> *Escherichia coli* strain BM2570

<400> 1711
atgaacaaaa atataaaaata ttctcaaaac tttttaacga gtgaaaaagt actcaaccaa 60
ataataaaaac aattgaattt aaaagaaacc gataccgttt acgaaattgg aacaggtaaa 120
gggcatttaa cgacgaaact ggctaaaata agtaaacagg taacgtctat tgaattagac 180

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agtcattctat tcaactttatc gtcagaaaaa ttaaaatcga atactcgtgt cacttttaatt 240
caccaagata ttctacagtt tcaattccct aacaaacaga ggtataaaaat tggtgggaat 300
attccttacc atttaagcac acaattattt aaaaaagtgg tttttgaaag ccatgcgtct 360
gacatctatc tgattgttga agaaggattc tacaagcgta ccttggtat taccgaaca 420
ctagggttgc tcttgccacac tcaagtctcg attcagcaat tgcttaagct gccagcggaa 480
tgctttcatc ctaaaccaag agtaaacagt gtcttaataa aacttaccg ccataccaca 540
gatgttccag ataaatattg gaagctatat acgtactttg tttcaaaatg ggtcaatcga 600
gaatatcgtc aactgtttac taaaaatcag tttcatcaag caatgaaaca cgccaaagta 660
aacaatttaa gtaccgttac ttatgagcaa gtattgtcta tttttaatag ttatctatta 720
tttaacggga ggaaataa

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<210> 1712
 <211> 735
 <212> DNA
 <213> *Staphylococcus aureus* strain RN451

```

<400> 1712
atgaacgaga aaaatataaa acacagtcaa aactttatta cttcaaaaaca taatatagat 60
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<210> 1713
 <211> 1029
 <212> DNA
 <213> *Enterococcus faecalis* strain V583

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<210> 1714
 <211> 818
 <212> DNA
 <213> *Campylobacter jejuni* subsp. *jejuni* ATCC 33292

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<400> 1714
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aacttgatat	cgtagataaa	agtggtgcgt	ggttttctta	taaagataaa	aaacttggac	780
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<210> 1715

<211> 809

<212> DNA

<213> *Abiotrophia adiacens* ATCC 49175

<400> 1715

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<210> 1716

<211> 817

<212> DNA

<213> *Abiotrophia defectiva* ATCC 49176

<400> 1716

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tgggtgttgt	actttaccag	aaggtactga	aatggttatg	ccaggcgaca	acgtacaaat	780
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<210> 1717

<211> 814

<212> DNA

<213> *Corynebacterium accolens* ATCC 49725

<400> 1717

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ccctcacacc	aagttcgagg	gttccgtcta	cgtcctgaag	aaggaagagg	gcggccgcca	660
caccccgyc	atgaacaact	accgtcctca	gttctacttc	cgcaccaccg	acgttaccgg	720
tggtgtgaac	ctgcctgagg	gcaccgagat	ggttatgcct	ggcgacaacg	ttgagatgtc	780
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<210> 1718

<211> 814

<212> DNA

<213> *Corynebacterium genitalium* ATCC 33031

<400> 1718

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<210> 1719

<211> 814

<212> DNA

<213> *Corynebacterium jeikeium* ATCC 43216

<400> 1719

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tggtgtgaag	ctgcctgagg	gcaccgagat	ggttatgccg	ggcgacaacg	tygacatgtc	780
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<210> 1720

<211> 748

<212> DNA

<213> *Corynebacterium pseudodiphtheriticum* ATCC 10700

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agaagtcgag	atcatcggca	tcaaggaaaa	gtcccagaag	accaccatca	ccggtatcga	480
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<210> 1721
<211> 813
<212> DNA
<213> *Corynebacterium striatum* ATCC 6940

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gttgacgacg	aggaaattat	cgagctcgtc	gagatggaga	tccgcgaact	gctcgcagag	180
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<212> DNA
<213> *Enterococcus avium* ATCC 14025

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<212> DNA
<213> *Gardnerella vaginalis* ATCC 14018

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 <212> DNA
 <213> *Listeria innocua* ATCC 33090

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<210> 1725
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 <212> DNA
 <213> *Listeria ivanovii* ATCC 19119

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ttccaactcc	agaacgtgat	actgacaaac	cattcatgat	gccagttgag	gatgtattct	360
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<210> 1726
 <211> 817
 <212> DNA
 <213> *Listeria monocytogenes* strain LSPQ 5093202

<400> 1726						
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<210> 1727

<211> 817

<212> DNA

<213> *Listeria seeligeri* ATCC 35967

<400> 1727

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<210> 1728

<211> 814

<212> DNA

<213> *Staphylococcus aureus* ATCC 25923

<400> 1728

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<210> 1729

<211> 817

<212> DNA

<213> *Staphylococcus saprophyticus* ATCC 15305

<400> 1729

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<210> 1730
 <211> 817
 <212> DNA
 <213> *Staphylococcus simulans* ATCC 27848

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<210> 1731
 <211> 817
 <212> DNA
 <213> *Streptococcus agalactiae* ATCC 27591

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<210> 1732
 <211> 817
 <212> DNA
 <213> *Streptococcus pneumoniae* ATCC 27336

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ggttgacgac	gaagaattgc	ttgaattggt	tgaaatggaa	atccgtgacc	tattgtcaga	180
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<210> 1733
 <211> 817
 <212> DNA
 <213> Streptococcus salivarius ATCC 7073

<400> 1733						
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<210> 1734
 <211> 897
 <212> DNA
 <213> Agrobacterium tumefaciens

<400> 1734						
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<210> 1735
 <211> 885
 <212> DNA
 <213> Bacillus subtilis strain 168

<400> 1735						
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<210> 1736

<211> 882

<212> DNA

<213> *Bacteroides fragilis* DSM 2151

<400> 1736

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<210> 1737

<211> 888

<212> DNA

<213> *Borrelia burgdorferi* strain U78183

<400> 1737

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<210> 1738

<211> 894

<212> DNA

<213> *Brevibacterium linens* DSM 20425

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<210> 1739

<211> 888

<212> DNA

<213> Chlamydia trachomatis strain F/IC-Cal-13

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<210> 1740

<211> 891

<212> DNA

<213> Fibrobacter succinogenes strain S85

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<400> 1740
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<210> 1741

<211> 894

<212> DNA

<213> *Flavobacterium ferrugineum* DSM 13524

<400> 1741

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<210> 1742

<211> 906

<212> DNA

<213> *Helicobacter pylori* strain 26695

<400> 1742

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<210> 1743

<211> 891

<212> DNA

<213> *Micrococcus luteus* strain IFO 3333

<400> 1743

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<210> 1744
<211> 891
<212> DNA
<213> *Mycobacterium tuberculosis* strain Erdmann

<400> 1744
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<210> 1745
<211> 891
<212> DNA
<213> *Mycoplasma genitalium* strain G37

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<210> 1746
<211> 891
<212> DNA
<213> *Neisseria gonorrhoeae* strain MS11

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<210> 1747
<211> 891
<212> DNA
<213> Rickettsia prowazekii strain Madrid E

<400> 1747
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<210> 1748
<211> 891
<212> DNA
<213> Salmonella choleraesuis subsp. choleraesuis strain LT2 trpE9

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<211> 881
<212> DNA
<213> Shewanella putrefaciens DSM 50426

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<210> 1750

<211> 897

<212> DNA

<213> *Stigmatella aurantiaca* strain DW4

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<210> 1751

<211> 897

<212> DNA

<213> *Thiomonas cuprina* strain Hoe5

<400> 1751

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<210> 1752

<211> 894

<212> DNA

<213> *Treponema pallidum* strain Nichols

<400> 1752

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cagttttatt ttagaactac tgacattacc ggtacgattt ctcttcctga aggggtagac 780
atgggtgaagc cgggggataa caccaagatt ataggtgagc tcatccacc gatagctatg 840
gacaagggtc tgaagcttgc gattcgtgaa ggggggcgca ctattgcttc tgggt 894

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<210> 1753
 <211> 891
 <212> DNA
 <213> *Ureaplasma urealyticum* ATCC 33697

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<400> 1753
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atcgttgttt tcttaaactaa atgtgatttc atgacagatc cagatatgca agatcttggt 180
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attcgtgggt caggtcttaa ggcttttaga ggagatccag tttgagaagc aaaaattgat 300
gaattaatgg acgcagttga ttcattgaatt ccattaccag aacgtagtac tgacaaacca 360
ttcttattag caattgaaga tgtattcaca atttcaggac gtggtacagt agtaactgga 420
cgtgttgaac gtggtgtatt aaaagttaat gatgaggttg aaattggttg tctaaagac 480
actcaaaaaa ctggtgttac aggaattgaa atggttagaa aatcattaga tcaagctgaa 540
gctggtgata atgctggtat tttattacgt ggtattaaaa aagaagatgt tgaacgtgg 600
caagtacttg taaaaccagg atcaattaaa cctcaccgta cttttactgc taaagtttat 660
attcttaaaa aagaagaagg tggacgtcat acacctattg tttcaggata ccgtccacaa 720
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ggtatgccag gtgatgacgt tgaaatgact gtgaattaa ttgctccagt tgcgattgaa 840
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<210> 1754
 <211> 909
 <212> DNA
 <213> *Wolinella succinogenes* DSM 1740

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<400> 1754
aacatgatta caggtgctgc tcaaattggat ggcgcgattc ttgttgtttc tgcggcgagat 60
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gaagatgttg agagaggtat ggttctttgt aaaataggtt ctatcactcc tcacactaac 660
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aatggatacc gacctcagtt ctatgttaga actacagacg ttaccggttc tatctctctt 780
cctgagggcg tagagatggt tatgcctggt gacaacgtta agatcaatgt tgagcttatc 840
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ggtgcgggt

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<210> 1755
 <211> 888
 <212> DNA
 <213> *Burkholderia cepacia*

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<400> 1755
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ccgatgccgc aaacgcgtga gcacatcctg ctggcgcgtc aggttggtgt tccgtacatc 120
atcgtgttcc tgaacaagtg cgacagtgtg gacgacgctg aactgctcga gctggctcag 180

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atggaagttc	gcgaaactcct	gtcgaagtac	gacttcccgg	gcgacgacac	gccgatcgtg	240
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atgagcctgg	cagacgcgct	ggacacgtac	atcccgcgc	cggagcgtgc	agttgacggc	360
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caggcaggcg	acaacgtcgg	tatcctgctg	cgcggcacga	agcgtgaaga	cgtggagcgt	600
ggccagggttc	tggcgaagcc	gggttcgatc	acgccgcaca	cgcacttcac	ggctgaagtg	660
tacgtgctga	gcaaggacga	aggcggccgt	cacacgccgt	tcttcaacaa	ctaccgtccg	720
cagttctact	tccgtacgac	ggacgtgacg	ggctcgatcg	agctgccgaa	ggacaaggaa	780
atggtgatgc	cgggcgacaa	cgtgtcgatc	acggtgaagc	tgattgctcc	gatcgcgatg	840
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<210> 1756

<211> 388

<212> DNA

<213> *Bacillus anthracis* strain CIP 9444

<400> 1756

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aggtgttaac	atcgatgaat	tactattatc	acaacctgat	acaggggagc	aaggtttaga	180
aatcgacagaa	gcacttgtag	gaagtgggtg	ggttgatatt	atcgtaattg	actctgtagc	240
agctcttgta	ccgaaagctg	aaattgaagg	agacatgggt	gactcacacg	taggtttaca	300
agctcgtcta	atgtctcaag	cacttcgtaa	actttcaggt	gcaatcaata	aatcaaaaaac	360
aatcgcaatc	tttattaacc	aaattcgt				388

<210> 1757

<211> 388

<212> DNA

<213> *Bacillus anthracis* ATCC 4229

<400> 1757

tgaaagtcca	ggtaaaacaa	cagtttcatt	acacgcaatt	gcagaagtac	agcgtcaagg	60
tggacaagca	gcgttcattg	atgctgagca	tgcaatggat	cctgtatatg	cacaaaaact	120
aggtgttaac	atcgatgaat	tactattatc	acaacctgat	acaggggagc	aaggtttaga	180
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agctcttgta	ccgaaagctg	aaattgaagg	agacatgggt	gactcacacg	taggtttaca	300
agctcgtcta	atgtctcaag	cacttcgtaa	actttcaggt	gcaatcaata	aatcaaaaaac	360
aatcgcaatc	tttattaacc	aaattcgt				388

<210> 1758

<211> 388

<212> DNA

<213> *Bacillus cereus* ATCC 7064

<400> 1758

tgaaagtcca	ggtaaaacaa	cagtttcatt	acacgcaatt	gcagaagtac	agcgtcaagg	60
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aggtgttaac	atcgatgaat	tactattatc	acaacctgat	acaggggagc	aaggtttaga	180
aatcgacagaa	gcacttgtag	gaagtgggtg	ggttgatatt	atcgtaattg	actctgtagc	240
agctcttgta	ccgaaagctg	aaattgaagg	agacatgggt	gactcacacg	taggtttaca	300
agctcgtcta	atgtctcaag	cacttcgtaa	actttcaggt	gcaatcaata	aatcaaaaaac	360
aatcgcaatc	tttattaacc	aaattcgt				388

<210> 1759

<211> 388

<212> DNA

<213> *Bacillus cereus* ATCC 13472

<400> 1759

tgaaagtcca	ggtaaaacga	cagtttcatt	acatgcaatt	gcagaagtac	aacgtcaagg	60
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tggacaagca gcattcatcg atgCGGagca cgcaatggat cctgtatatg cacaaaaatt 120
aggcgTTaAc atagatgaat tactattatc acagcctgat acaggggagc aaggattaga 180
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agcacgtTTa atgtcacaag cacttcgTaa gctttcagga gcaatcaaca aatcaaaaac 360
aattgcaatc tttattaacc aaattcgt 388
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<210> 1760

<211> 374

<212> DNA

<213> *Bacillus mycoides* ATCC 6462

<400> 1760

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tcatcgatgc ggagcacgca atggatcctg tatatgcaca aaaattaggc gttaacatag 120
atgaattact attatcacag cctgatacag gggagcaagg attagaaatc gcagaagcac 180
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aagcagagat tgaaggagac atgggtgact cacacgtagg tttacaagca cgtttaatgt 300
cacaagcact tcgtaagctt tcaggagcaa tcaacaaatc aaaaacaatt gcaatcttta 360
ttaaccaaTt tcgt 374
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<210> 1761

<211> 381

<212> DNA

<213> *Bacillus pseudomyoides* NRRL BD-10

<400> 1761

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ggaaagtTca ggtaaaacaa cggtttcctt acatgcgatt gcagaagtgc aacgtcaagg 60
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aggTgttaat attgatgagt tactattatc gcagcctgat acaggagaac aaggTttaga 180
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agctcttTgta ccaaaagcag aaatcgaaagg ggaatgggt gactccacg ttggTttaca 300
agcgcgTtTa atgtcacaag cacttcgTaa gctttctggT gcgattaaca aatcaaaaac 360
aattgcaatc ttcattaacc a 381
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<210> 1762

<211> 388

<212> DNA

<213> *Bacillus thuringiensis* strain HER 1410

<400> 1762

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aggcgTTaAc atagatgaat tactattatc acagcctgat acaggggagc aaggattaga 180
aatcgCGgaa gcacttgtac gaagtggTgc ggttgacatt atcgtaattg actctgtagc 240
agctcttTgta ccgaaagcag agattgaagg cgacatgggt gactcacacg taggtttaca 300
agcacgtTTa atgtcacaag cacttcgTaa gctttcagga gcaatcaaca aatcaaaaac 360
aattgcaatc tttattaacc aaattcgt 388
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<210> 1763

<211> 388

<212> DNA

<213> *Bacillus thuringiensis* strain HER 1418

<400> 1763

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aggcgTTaAc atagatgaat tactattatc acagcctgat acaggggagc aaggattgga 180
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agcacgtTTa atgtcacaag cacttcgTaa gctttcagga gcaatcaaca aatcaaaaac 360
aattgcaatc tttattaacc aaattcgt 388
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<210> 1764
<211> 358
<212> DNA
<213> *Klebsiella oxytoca* ATCC 33496

<400> 1764
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gaacaaagcc tataaaaaat ctgcccgtgt cgtgggtgac gtcacggta aataccacc 180
tcatggtgat actgccgtat acgacaccat tgtacgtatg gcgcagccct tctccctgcg 240
ttacatgctg gtagatggcc agggtaactt tggttcggtc gacggcgact ccgccgcagc 300
gatgcgttat acggaaatcc gtatgtcgaa gatcgcccat gaactgatgg ccgacctc 358

<210> 1765
<211> 365
<212> DNA
<213> *Klebsiella pneumoniae* subsp. *ozaenae* ATCC 11296

<400> 1765
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tccgagatgg cctgaagccg gtacaccgtc gcgtacttta cgccatgaac gtattgggca 120
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ccgcggcgat gcgttatacc gaaattcgtc tggcgaaaat cgctcatgag ctgatggccg 360
atctt 365

<210> 1766
<211> 344
<212> DNA
<213> *Klebsiella planticola* ATCC 33531

<400> 1766
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cgagatggcc tgaaacccgt acaccgtcgc gtactttacg ccatgaacgt attgggcaat 120
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caccctcatg gtgataccgc cgtttatgac accattgtac gtatggcaca gccattctcc 240
ttgcgttata tgctggtcga tggccagggt aacttcgggt ctgtcgatgg cgactccgcc 300
gcagcgatgc gttatacggg aatccgtatg tcgaaaatcg ccca 344

<210> 1767
<211> 345
<212> DNA
<213> *Klebsiella pneumoniae* ATCC 27336

<400> 1767
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gatggcctga agccggtaca ccgtcgcgta ctttacgcca tgaacgtatt gggcaatgac 120
tggaaacaaag cctataaaaa atcagcccgt gtcgttgggt acgtaatcgg taaataccac 180
ccgcacggcg actccgcggg atacgacacc atcgtgcgta tggcgagcc gttctcgctg 240
cgttacatgc tgggtggacgg ccagggtaac tttggttcca tcgacggcga ctccgccgcg 300
gcgatgcggt ataccgaaat tcgtctggcg aaaatcgctc atgag 345

<210> 1768
<211> 356
<212> DNA
<213> *Klebsiella pneumoniae* subsp. *pneumoniae* ATCC 13883

<400> 1768
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gcctgaagcc ggtacaccgt cgcgtacttt acgcatgaa cgtattgggc aatgactgga 120

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acatgctggt ggacggccag ggtaactttg gttccatcga cggcgactcc gccgcggcga 300
tgcgttatac cgaaattcgt ctggcgaaaa tcgctcatga gctgatggcc gatctt 356
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<210> 1769

<211> 361

<212> DNA

<213> *Klebsiella pneumoniae* subsp. *pneumoniae* ATCC 29011

<400> 1769

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ccgagatggc ctgaagccgg tacaccgtcg cgtactttac gccatgaacg tattgggcaa 120
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gctgcgttac atgctggtgg acggccaggg taactttggt tccatcgacg gcgactccgc 300
cgcgccgatg cgttataccg aaattcgtct ggcgaaaatc gtcctatgagc tgatggccga 360
t 361
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<210> 1770

<211> 365

<212> DNA

<213> *Klebsiella pneumoniae* subsp. *rhinoscleromatis* ATCC 13824

<400> 1770

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cgcgccgatg cgttataccg aaattcgtct ggcgaaaatc gtcctatgagc tgatggccga 360
tcttg 365
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<210> 1771

<211> 357

<212> DNA

<213> *Klebsiella terrigena* ATCC 33257

<400> 1771

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tatatgctgg tcgatggcca gggtaacttc ggttctgtcg atggcgactc cgccgcagcg 300
atgcgttata cggaaatccg tatgtcgaaa atcgccacg agctgatggc cgacctc 357
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<210> 1772

<211> 968

<212> DNA

<213> *Legionella pneumophila* strain *pneumophila* ATCC 33152

<400> 1772

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<210> 1773

<211> 967

<212> DNA

<213> *Proteus mirabilis* ATCC 25933

<400> 1773

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<210> 1774

<211> 978

<212> DNA

<213> *Providencia rettgeri* ATCC 9250

<400> 1774

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<210> 1775

<211> 978

<212> DNA

<213> *Proteus vulgaris* ATCC 13315

<400> 1775

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<210> 1776

<211> 940

<212> DNA

<213> *Yersinia enterocolitica* ATCC 9610

<400> 1776

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<210> 1777

<211> 668

<212> DNA

<213> *Klebsiella oxytoca* ATCC 13182

<400> 1777

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<210> 1778

<211> 714

<212> DNA

<213> *Klebsiella oxytoca* ATCC 33496

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ccagaatggc cgcgggttcgg tgcggatgcg cgcggtatgg gccaaagaag acggcgcggg 660
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<210> 1779

<211> 722

<212> DNA

<213> Klebsiella pneumoniae subsp. ozaenae ATCC 11296

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<210> 1780

<211> 692

<212> DNA

<213> Klebsiella planticola ATCC 33531

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<210> 1781

<211> 700

<212> DNA

<213> Klebsiella pneumoniae ATCC 27336

<400> 1781
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<210> 1782

<211> 726

<212> DNA

<213> *Klebsiella pneumoniae* subsp. *pneumoniae* ATCC 13883

<400> 1782

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<210> 1783

<211> 706

<212> DNA

<213> *Klebsiella pneumoniae* subsp. *pneumoniae* ATCC 29011

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<210> 1784

<211> 614

<212> DNA

<213> *Klebsiella pneumoniae* subsp. *rhinoscleromatis* ATCC 13884

<400> 1784

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acgggcgcgg ctca 614

<210> 1785
<211> 668
<212> DNA
<213> *Klebsiella terrigena* ATCC 33257

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tgccgcac 668

<210> 1786
<211> 113
<212> DNA
<213> *Bacillus cereus* ATCC 7064

<400> 1786
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<210> 1787
<211> 118
<212> DNA
<213> *Bacillus cereus* ATCC 14579

<400> 1787
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<210> 1788
<211> 120
<212> DNA
<213> *Bacillus anthracis* strain CIP 9444

<400> 1788
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<210> 1789
<211> 118
<212> DNA
<213> *Bacillus cereus* ATCC 13472

<400> 1789
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tgaagaagta ccaaagtctg tttctgaaga aattatcaaa aaaaataaag gtgaataa 118

<210> 1790
<211> 120

<212> DNA

<213> *Bacillus anthracis* ATCC 4229

<400> 1790

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tatgaagaag taccaaagtc tgtttctgaa gaaattatca aaaaaataa aggtgaataa 120

<210> 1791

<211> 120

<212> DNA

<213> *Bacillus pseudomycoides* NRRL B-617

<400> 1791

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<210> 1792

<211> 116

<212> DNA

<213> *Bacillus cereus* ATCC 49064

<400> 1792

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<210> 1793

<211> 120

<212> DNA

<213> *Bacillus anthracis* strain CIP 9440

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<210> 1794

<211> 100

<212> DNA

<213> *Bacillus cereus* ATCC 15816

<400> 1794

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tgtttctgaa gaaattatca aaaaaataa aggtgaataa 100

<210> 1795

<211> 120

<212> DNA

<213> *Bacillus weihenstephanensis* strain WSBC 10204

<400> 1795

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<210> 1796

<211> 120

<212> DNA

<213> *Bacillus mycoides* ATCC 6462

<400> 1796

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<210> 1797
<211> 120
<212> DNA
<213> *Bacillus thuringiensis* ATCC 10792

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<210> 1798
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<212> DNA
<213> *Bacillus weihenstephanensis* strain WSBC 10204

<400> 1798
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gtttcacttt ctagtctaaa tataaaataa cccatataaa ctaaggagga atttaga 117

<210> 1799
<211> 117
<212> DNA
<213> *Bacillus thuringiensis* ATCC 10792

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<210> 1800
<211> 117
<212> DNA
<213> *Bacillus anthracis* ATCC 4229

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<210> 1801
<211> 117
<212> DNA
<213> *Bacillus pseudomycoides* NRRL B-617

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<210> 1802
<211> 117
<212> DNA
<213> *Bacillus anthracis* strain CIP 9444

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<210> 1803
<211> 117
<212> DNA
<213> *Bacillus cereus* ATCC 7064

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<210> 1804
<211> 117
<212> DNA
<213> *Bacillus cereus* ATCC 49064

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<210> 1805
<211> 117
<212> DNA
<213> *Bacillus mycoides* ATCC 6462

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<210> 1806
<211> 117
<212> DNA
<213> *Bacillus cereus* ATCC 14579

<400> 1806
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gtttcacttt ctagtctaaa tataaaataa cctatataaa ctaaggagga atttaga 117

<210> 1807
<211> 117
<212> DNA
<213> *Bacillus cereus* ATCC 15816

<400> 1807
ttgatttttta tgcgattcttc acgtataact acttatgtaa gcttagaaaag tgggacgcaa 60
gtttcgcttt ctagcctaaa tataaaataa cctatataaa ctaaggagga atttaga 117

<210> 1808
<211> 117
<212> DNA
<213> *Bacillus cereus* ATCC 13472

<400> 1808
ttgatttttta tgcgattgttc aagtataact acttatgtaa gcttagaaaag tgggacgtaa 60
gtttcacttt ctagtctaaa tataaaataa cctatataaa ctaaggagga atttaga 117

<210> 1809
<211> 116
<212> DNA
<213> *Bacillus anthracis* strain CIP 9440

<400> 1809
ttgatttttta tgcgattcttc acgtataact acttatgtaa gcttagaaaag tgggacgcaa 60
gtttcgcttt ctagcctaaa tataaaataa cctatataaa ctaaggagga atttag 116

<210> 1810

<211> 278
<212> DNA
<213> *Bacillus mycoides* ATCC 6462

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<400> 1810
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gttgaccatg gtataaactac attaaactgct gcgatcacta cagttcttgc aaaagctggg 120
ggtgctgaag cagcgggata cgatcaaatac gacgtgctc cagaagaaag agagcgcgga 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaaa aacatgat 278
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<210> 1811
<211> 278
<212> DNA
<213> *Bacillus thuringiensis* ATCC 10792

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<400> 1811
atgggctaaag ctaaattcga acgttctaaa ccccatgtta acatcgggtac aatcgggccac 60
gttgaccatg gtataaactac attaaactgct gcgatcacta cagttcttgc aaaagctggg 120
ggtgctgaag cagcgggata cgatcaaatac gatgctgctc cagaagaaag agagcgcggt 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaaa aacatgat 278
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<210> 1812
<211> 270
<212> DNA
<213> *Bacillus cereus* ATCC 15816

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<400> 1812
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gttgaccatg gtataaactac attaaactgct gcgatcacta cagtacttgc aaaagctggg 120
ggtgctgaag cagcgggata cgatcaaatac gatgctgctc cagaagaaag agagcgcggt 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaaa 270
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<210> 1813
<211> 278
<212> DNA
<213> *Bacillus weihenstephanensis* strain WSBC 10204

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atgggctaaag ctaaattcga acgttctaaa ccccatgtta acatcgggtac aatcgggccac 60
gttgaccatg gtataaactac attaaactgct gcgatcacta cagttcttgc aaaagctggg 120
ggtgctgaag cagcgggata cgatcaaatac gacgtgctc cagaagaaag agagcgcgga 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaaa aacatgat 278
```

<210> 1814
<211> 266
<212> DNA
<213> *Bacillus anthracis* strain CIP 9440

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<400> 1814
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ggtgctgaag cagcgggata cgatcaaatac gatgctgctc cagaagaaag agagcgcggt 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgt 266
```

<210> 1815
<211> 269

<212> DNA

<213> *Bacillus cereus* ATCC 7064

<400> 1815

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ggtgctgaag cagcgggata cgatcaaatac gatgctgctc cagaagaaaag agagcgcggt 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaa                269
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<210> 1816

<211> 268

<212> DNA

<213> *Bacillus cereus* ATCC 13472

<400> 1816

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atgggctaaag ctaaattcga acgttctaaa ccccatgtta acatcggtac aatcggccac 60
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ggtgctgaag cagcgggata cgatcaaatac gatgctgctc cagaagaaaag agagcgcggt 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaa                268
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<210> 1817

<211> 278

<212> DNA

<213> *Bacillus anthracis* ATCC 4229

<400> 1817

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atgggctaaag ctaaattcga acgttctaaa ccccatgtta acatcggtac aatcggccac 60
gttgaccatg gtaaaactac attaaactgct gcgatcacta cagtacttgc aaaagctggg 120
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gactgcccag gtcacgctga ctatgttaaa aacatgat                278
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<210> 1818

<211> 268

<212> DNA

<213> *Bacillus cereus* ATCC 14579

<400> 1818

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ggtgctgaag cagcgggata cgatcaaatac gatgctgctc cagaagaaaag agagcgcggt 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaa                268
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<210> 1819

<211> 278

<212> DNA

<213> *Bacillus anthracis* strain CIP 9444

<400> 1819

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atgggctaaag ctaaattcga acgttctaaa ccccatgtta acatcggtac aatcggccac 60
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ggtgctgaag cagcgggata cgatcaaatac gatgctgctc cagaagaaaag agagcgcggt 180
atcacaatct caactgcaca cgttgagtag gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaaa aacatgat                278
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<210> 1820

<211> 278

<212> DNA

<213> *Bacillus pseudomycoides* NRRL B-617

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<400> 1820
atggcctaaag ctaaattcga acgttctaaa ccccatgtta acatcggtac aatcggccac 60
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ggtgctgaag cagcgcgata cgaccaaatac gatgctgctc cagaagaaaag agagcgcggg 180
atcacaaatct caactgcaca cgttgagtac gaaactgaaa ctcgtcacta tgcacacggt 240
gactgcccag gtcacgctga ctatgttaaa aacatgat
278
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<210> 1821

<211> 263

<212> DNA

<213> *Bacillus cereus* ATCC 49064

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<400> 1821
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ggtgctgaag cagcgcgata cgatcaaatac gatgctgctc cagaagaaaag agagcgcggg 180
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263
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<210> 1822

<211> 1668

<212> DNA

<213> *Streptococcus oralis* ATCC 35037

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cacgtatcgt atttgctaac aaaatggaca aaatcggtgc tgacttcctt tactcagtaa 180
gcacacttca cgaccgtctt caagcaaacg cacacccaat ccaattgcc aatcggtgctg 240
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1668
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<210> 1823

<211> 115

<212> DNA

<213> *Budvicia aquatica* ATCC 35567

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<400> 1823
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cgaagcgcca aacaacgttg ctacagcaat cattgaagct cgtaaggcta gataa 115

<210> 1824

<211> 107

<212> DNA

<213> *Buttiauxella agrestis* ATCC 33320

<400> 1824

ctgcgttcac tgaccaaggt cgtgcatctt actccatgga attcctgaag tatgatgacg 60
cgccaaacaa cgtagctcag gccgtaatcg aagctcgcgg taaataa 107

<210> 1825

<211> 79

<212> DNA

<213> *Klebsiella oxytoca* ATCC 13182

<400> 1825

ttactccatg gagttcctga agtatgatga tgcgccgaac aacgttgctc aggccgtaat 60
cgaagcccggt ggtaaataa 79

<210> 1826

<211> 111

<212> DNA

<213> *Plesiomonas shigelloides* ATCC 14029

<400> 1826

cagctgcggt ctctgaccaa aggtcgtgca tcatacacta tggaaattcct gaagtatgat 60
gatgcgccaa acaacgttgc tcaggccggt attgaagccc gtggtaagta a 111

<210> 1827

<211> 108

<212> DNA

<213> *Shewanella putrefaciens* ATCC 8071

<400> 1827

gatttgcgct ctgcaactca tgggcgtgct tcgtactcca tggagttcct gaagtactct 60
gatgcaccgc aaaacattgc gaaagcgatt attgaatctc gtagctaa 108

<210> 1828

<211> 113

<212> DNA

<213> *Obesumbacterium proteus* ATCC 12841

<400> 1828

ctcagctgcg ttctctgacc aaaggctcgtg catcttactc catggaattc ctgaagtatg 60
atgatgcgcc taacaacgtt gctcaggccg ttattgaagc tcgtggcaaa taa 113

<210> 1829

<211> 70

<212> DNA

<213> *Klebsiella oxytoca* ATCC 13182

<400> 1829

gccgcagggt taaaacacaaa gtcccgtgct ctctcctgaa ggggagagca ctatagtaag 60
gaatatagcc 70

<210> 1830

<211> 66

<212> DNA

<213> *Budvicia aquatica* ATCC 35567

<400> 1830
gcctcgggta aaacttatat cccagtcgcc ctcgtataga gggggataga gtaaaggaag 60
ataatc 66

<210> 1831

<211> 81

<212> DNA

<213> *Plesiomonas shigelloides* ATCC 14029

<400> 1831
tccacaggat taaaacccag gtttaaacct aagtcgccgtg ctctctctc aggggagagc 60
acaatagtaa ggaatatagc c 81

<210> 1832

<211> 70

<212> DNA

<213> *Obesumbacterium proteus* ATCC 12841

<400> 1832
gctactagtt taaaacattg atccccgtgct ctctctatga agggagagca caagagtaag 60
gaataagcc 70

<210> 1833

<211> 72

<212> DNA

<213> *Shewanella putrefaciens* ATCC 8071

<400> 1833
tttccagtta cgacataaat gttattatgg tccagctttg actggactat tctgaaaaga 60
aaggaatata tc 72

<210> 1834

<211> 73

<212> DNA

<213> *Buttiauxella agrestis* ATCC 33320

<400> 1834
gccccgggtt ttaaaaaaca ttgatccccgt gctctctcca gaaggggaga ggcgaacagt 60
aaggaatata gcc 73

<210> 1835

<211> 795

<212> DNA

<213> *Campylobacter coli* ATCC 43479

<400> 1835
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caaaaaaaga agaagttatc cgcggtatgg ttcttgctaa accaaaatca attactccac 600
atactgattt cgaagctgaa gtttatatcc taaataaaga tgaggggtgg agacatactc 660
cattctttta taactataga ccgcaattct atgtaagaac aacagatgta acaggttcta 720
ttaaattagc tgatggcggt gaaatgggtta tgcctgggtga aaatgtaaga attactgtaa 780

gcttgattgc accag

795

<210> 1836

<211> 817

<212> DNA

<213> *Campylobacter fetus* subsp. *fetus* ATCC 25936

<400> 1836

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ctatctcgtc	aagttggtgt	tccatatata	gttgttttta	tgaacaaagc	tgatatggta	120
gatgacgcag	agttgctaga	attagttgaa	atggagatca	gagagttatt	aagcgaatat	180
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gttaaagttg	gtgatactat	cgaaatcgta	ggtattagag	atacacaaac	tacaacagtt	480
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gttcttttac	gcggtacaaa	gaaagaagac	gttgaaagag	gtatggttct	ttgtaagcca	600
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ggcggtagac	atactccatt	cttcaacaac	tatagaccac	aattttatgt	aagaacaaca	720
gatgttactg	gatcaatcac	tcttccagag	ggtactgaga	tggttatgcc	tggtgataac	780
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<210> 1837

<211> 798

<212> DNA

<213> *Campylobacter fetus* subsp. *venerealis* ATCC 33561

<400> 1837

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aatcaattac	tcctcatact	aaatttgagg	gagaagttta	tatcttgact	aaggaagagg	660
gcggtagaca	tactccattc	ttcaacaact	atagaccaca	attttatgta	agaacaacag	720
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taaaaatcac	tgtagtt					798

<210> 1838

<211> 1116

<212> DNA

<213> *Buttiauxella agrestis* ATCC 33320

<400> 1838

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ccagaaccag	agcgtgctat	cgacaagcca	ttcctgctgc	caatcgaaga	cgtattctcc	660
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actccgttct	tcaaaggcta	ccgtccacag	ttctacttcc	gtacaactga	cgtgactggc	1020
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<210> 1839

<211> 1109

<212> DNA

<213> *Klebsiella oxytoca* ATCC 13182

<400> 1839

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<210> 1840

<211> 1108

<212> DNA

<213> *Plesiomonas shigelloides* ATCC 14029

<400> 1840

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<210> 1841

<211> 1107

<212> DNA

<213> *Shewanella putrefaciens* ATCC 8071

<400> 1841

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<210> 1842

<211> 1116

<212> DNA

<213> *Obesumbacterium proteus* ATCC 12841

<400> 1842

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ggtgatgatg	aagagctgct	ggagctggta	gaaatggaag	ttcgtgaact	tctgtctcag	480
tacgacttcc	caggcaatga	tactccaatc	atccgtgggt	ctgctctgaa	agcgctggaa	540
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<210> 1843

<211> 1129

<212> DNA

<213> *Budvicia aquatica* ATCC 35567

<400> 1843

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<210> 1844

<211> 810

<212> DNA

<213> *Abiotrophia adiacens* ATCC 49175

<400> 1844

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<210> 1845

<211> 815

<212> DNA

<213> *Arcanobacterium haemolyticum* ATCC 9345

<400> 1845

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ccaggcaggt	tcggaagtgt	ccacgttgct	tggccgtatg	ccatcagcag	tgggctacca	780
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<210> 1846

<211> 1073

<212> DNA

<213> *Basidiobolus ranarum* ATCC 24670

<400> 1846

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ccacccccga	aatcttggag	actggtatta	aggttgtega	tttgttggcc	ccttacgctc	360

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<210> 1847

<211> 480

<212> DNA

<213> *Blastomyces dermatitidis* ATCC 56220

<400> 1847

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gtgtctgccc	ttttgggtcg	tatccccctt	gccgtcgggt	accagcccac	tctcgcctgc	420
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<210> 1848

<211> 566

<212> DNA

<213> *Blastomyces dermatitidis* ATCC 14112

<400> 1848

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<210> 1849

<211> 817

<212> DNA

<213> *Campylobacter coli* ATCC 43479

<400> 1849

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<210> 1850

<211> 775

<212> DNA

<213> *Campylobacter fetus* subsp. *fetus* ATCC 25936

<400> 1850

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<210> 1851

<211> 793

<212> DNA

<213> *Campylobacter fetus* subsp. *venerealis* ATCC 33561

<400> 1851

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<210> 1852

<211> 825

<212> DNA

<213> *Campylobacter gracilis* ATCC 33236

<400> 1852

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gtccgcgctt ttaggacgaa ttccgtccgc ggtcggttat cagcctacgc ttgccagcga 780

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825

<210> 1853

<211> 818

<212> DNA

<213> *Campylobacter jejuni* subsp. *jejuni* ATCC 33560

<400> 1853

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tatgacagat	ggtttggtta	ggggcttaaa	agctgaggct	ttaggtgctc	ctattagtgt	180
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tgtgcttatg	tttattgata	atatctttag	attttcacaa	tcaggttctg	aaatgtcagc	720
acttttagga	agaattccat	cagctgtggg	ttatcaacca	accctagcaa	gtgaaatggg	780
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<210> 1854

<211> 830

<212> DNA

<213> *Enterococcus cecorum* ATCC 43198

<400> 1854

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aacggatggt	ttacaacgtg	ggatggaagt	tgctcgatact	ggtaaaccaa	tttcagttcc	180
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tgtgttgcta	tttattgata	acatcttccg	tttcaactca	gcgggttctg	aagtatcagc	720
cttgcttggt	cgtatgccat	ctgccgtggg	ttatcaacct	acattggcta	cagaaatggg	780
tcaattacaa	gaacgtatca	cttcaactaa	gaagggctct	atcacttcta		830

<210> 1855

<211> 823

<212> DNA

<213> *Enterococcus columbae* ATCC 51263

<400> 1855

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atcgtgcttg	aagctgcttt	agagctagga	gatggcatta	ttcgtacgat	tgcaatggaa	120
tcaactgatg	gattgcaacg	tggaatggaa	gttttcgata	caggtaagcc	aatttcagta	180
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caagaagctt	ttcctgctga	tgcgaatcgt	gatgcgattc	ataaatcagc	tcagctttt	300
gaagaattaa	gtacaagtac	tgaaatccta	gaaacaggga	ttaaagttat	cgacttacta	360
gcaccatact	taaaaggtgg	gaaagttggt	ctattcgggtg	gtgccgggtg	aggtaaaacc	420
gtattaattc	aagaattaat	tcataatatc	gcccaagaac	atgggggtat	ttcagtatct	480
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ggcgttattc	aaaaaactgc	tatgggtgtt	gggcaaatga	acgaaccacc	tggagcacgt	600
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gacgtattgc	tattttattga	taatatcttc	cgttttactc	aagcagggtc	tgaagtttct	720
gccttacttg	gtcgtatgcc	ttctgcggta	ggttatcaac	ctactttggc	tactgaaatg	780

ggtcaattgc aagaacggat tacatcaacg aagaaagggt cga

823

<210> 1856

<211> 826

<212> DNA

<213> *Enterococcus dispar* ATCC 51266

<400> 1856

ttaccagaca	ttaataatgc	cttggttgtc	tataaaaaatg	acgaacaaaa	aaccaagatt	60
gtattagaag	ctgccttaga	actaggagat	ggtgtgattc	gaactatcgc	catggaatct	120
actgatggct	tacaacgggg	aatggaagtt	gtcgatactg	gcagttccat	ttctgtaccg	180
gtaggaaaag	aaacattggg	tcgtgtatct	aacgttttag	gaaatacaat	tgacttagaa	240
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gaattaagca	ctagtacaga	aatttttagaa	acagggatta	aagttattga	cctattagcc	360
ccttatttaa	aaggtggtaa	agtcggatta	ttcggtggtg	cggagttgg	taaaaccggt	420
ttaattcaag	aattaattca	taatatggcc	caagaacatg	gtgggatttc	tgtttttact	480
ggtgttggtg	aaagaacacg	tgaaggtaat	gacttgtatt	atgaaatgaa	agaatctggc	540
gttatcgaaa	aaactgccat	ggtatttggt	caaatgaatg	agccacctgg	tgcccggatg	600
cgggttgctt	taaccggact	taccattgcg	gaatacttcc	gggacgttga	aggacaagat	660
gtattgctct	ttatcgataa	tattttccgt	tttacccaag	ctgggttcaga	agtatctgcc	720
ttattaggac	ggatgccctc	tgccgttggt	tatcaaccaa	ctttggctac	tgaaatggga	780
caacttcaag	aacggattac	ctcaacgaaa	aaaggttcta	ttacat		826

<210> 1857

<211> 814

<212> DNA

<213> *Enterococcus malodoratus* ATCC 43197

<400> 1857

tccttaccag	acatcaacaa	tgcgttgatt	gtttacaaaa	aaaataaaac	aaaagttggt	60
cttgaagctg	ctttggaact	tggtgatggt	gttatccgca	cgatctctat	ggaatcaaca	120
gatggcttgc	aacgtggaat	ggaagttgtc	gatacaggca	aaccaatctc	agttcccgtt	180
ggtaaaagaa	ctttaggctg	tgtgtttaac	gtattagggt	aaacaatcga	caaagaagcg	240
ccttttccag	aagatgcagt	aaaaagcggg	attcataaaa	aagcgccggc	ttttgaagaa	300
cttagtacca	gtaatgaaat	tttagaaaca	gggatcaaag	ttatcgactt	attagctcct	360
tacttaaagg	gtggtaaagt	cggactatct	ggtggtgccc	gtgttggtta	aaccgtcttg	420
atccaagaat	tgattcataa	tatcgcccaa	gaacacgggt	gtatttcagt	gtttacgggt	480
ggtgtgtaac	gtactcgtga	agggaaacgac	ctttattatg	aaatgaagga	atcaggcgtt	540
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ctgttggtta	tcgacaacat	cttccgtttc	actcaagccg	gttctgaagt	ttctgccttg	720
cttggccgga	tgctttcagc	cgttggtctac	caaccaactt	tggcaactga	aatgggtcaa	780
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<210> 1858

<211> 791

<212> DNA

<213> *Enterococcus mundtii* ATCC 43186

<400> 1858

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agaattaggt	gacggtgtga	tccgtacgat	cgcaatggaa	tcgacggatg	gactacaacg	120
tggaatggaa	gtcatcgaca	caagcaaagc	gatctctgta	ccagttggaa	cagaaacatt	180
aggctcgtgtg	ttcaacgtgt	taggtgaaac	aatcgatttg	gaagcaccat	ttccagagga	240
tgcccaaaga	agcgagatcc	acaagaaagc	accaaatttt	gatgaattaa	gcacaagtag	300
agagattcct	gaaactggga	tcaaagtcac	tgacttatta	gcaccttatt	taaaagggtg	360
gaaagttgga	ttgtttgggg	gtgccgggtg	tggtaaaacc	gtactgatcc	aagaattgat	420
ccataatatc	gcccaagaac	atgggggaat	ctcagtgttt	accggtgtag	gggaacgtac	480
ccgtgaagga	aacgatctgt	attacgaaat	gaaagattca	ggcgtaaatcg	aaaaaacagc	540
gatggtgttt	ggacaaatga	atgagccacc	aggtgctcgt	atgcgtgtcg	cactaactgg	600
attgacgatt	cgggaatatt	tccgtgatgt	cgaaggacaa	gacgtgctct	tattttattga	660
taatattttc	cgtttcacc	aagcaggttc	agaagtatct	gccttactag	gacgtatgcc	720
atcagcgggt	ggttatcaac	caacccttagc	gactgaaatg	ggacaactcc	aagaacggat	780

cacttcaacg a

791

<210> 1859

<211> 817

<212> DNA

<213> *Enterococcus raffinosus* ATCC 49427

<400> 1859

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gatggattac	aacgtggaat	ggaagttgtc	gatactggca	agcctatttc	tgttccagta	180
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cttggccgta	tgccgtcagc	agttggttac	caaccgactt	tagcaactga	aatgggtcaa	780
ttacaagaac	gtattacgtc	gacgaaaaaa	ggttcaa			817

<210> 1860

<211> 852

<212> DNA

<213> *Globicatella sanguis* ATCC 51173

<400> 1860

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atggtccgga	caattgcat	ggaatcaacc	gatggtttgg	aacgcggcat	gacagttgtg	180
gattatttaa	caccgattaa	agtgccagta	ggcgaagcca	ctttaggtag	agtattcaat	240
gttttaggtg	agacaattga	tgaactagaa	ccggttggcg	acgatgttga	actcaaaagt	300
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ctcgtttttg	aaatgcgaga	gtcagggtga	agcaagaaga	cggccatggt	tttcggtcaa	600
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caaccaactt	tagcaagtga	aatgggacaa	atgcaagaac	gtattacgtc	wacgaagcgc	840
ggttccatta	ca					852

<210> 1861

<211> 828

<212> DNA

<213> *Lactococcus garvieae* ATCC 49156

<400> 1861

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gagcacgtat	gcgtgttgct	cttactgggt	tgacaattgc	tgaatatttc	cgtgatgtag	660
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aagtatctgc	cctcttagga	cgtatgccat	cagccgttgg	ttaccaacct	acgcttgcaa	780
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<210> 1862
 <211> 828
 <212> DNA
 <213> *Lactococcus lactis* ATCC 11454

<400> 1862						
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<210> 1863
 <211> 825
 <212> DNA
 <213> *Listeria ivanovii* ATCC 19119

<400> 1863						
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<210> 1864
 <211> 821
 <212> DNA
 <213> *Succinivibrio dextrinosolvens* ATCC 19716

<400> 1864						
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<210> 1865

<211> 822

<212> DNA

<213> *Tetragenococcus halophilus* ATCC 33315

<400> 1865

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gtttaatgg	gtagggtgaac	gtactcgtga	aggtaatgac	ttgtattatg	aaatgcagga	540
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tcaagacgta	ttattattta	ttgataatat	tttccgtttt	acacaagcag	gtaccgaagt	720
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aatggggcaa	ctgcaagaac	ggattacgtc	aacggataag	gg		822

<210> 1866

<211> 818

<212> DNA

<213> *Campylobacter fetus* subsp. *fetus* ATCC 25936

<400> 1866

atctcctcag	gatckatagg	acttgatata	gctcttggtg	taggcggcgt	acaaaaagga	60
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atagcagaat	ctcaaaaagt	cggcggagtt	tgcgcgtttg	tagatgcaga	gcatgcactt	180
gatgttaaat	atgctaaaaa	tttaggcgtt	gatacggata	acttatatat	ttctcaaccg	240
gacttcggag	agcaagctct	tgatatagta	gaaactctag	ctagaagcgg	cgccgttgat	300
cttatagtaa	tagatagcgt	agcagctyta	acaccaaaaa	gcgaaataga	aggcgatatg	360
ggagatcagc	acgtagggct	gcaagcaaga	ctcatgagtc	aagcacttag	aaaattaac	420
ggagttgtcc	ataaaatggg	aactacagtt	gtattttata	accaaattcg	tatgaaaatc	480
ggcgctatgg	gctatggcac	tcctgaaact	actactggcg	gaaatgcgct	taaattttac	540
gcttcagtta	gacttgacgt	acgtaaaata	gctactttta	aacagagcga	tgagccaatc	600
ggaaaccgcg	taaaagtaaa	agtagtaaaa	aacaaagtcg	ctcctccttt	tagacaagcc	660
gaatttgata	tcatgttttg	agaaggtatc	agcaagaag	gagagataat	agattacggc	720
gtaaaacttg	atattatcga	taaaagcggc	gcttggttta	gctatgataa	ttcaaaaatta	780
ggtcaaggca	gagaaaactc	aaaagcgttt	ttaaaaga			818

<210> 1867

<211> 814

<212> DNA

<213> *Campylobacter fetus* subsp. *venerealis* ATCC 33561

<400> 1867

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tagcagaatc	tcaaaaagtc	ggcggagttt	gcgcgtttgt	agatgcagag	catgcacttg	180
atgttaaaat	tgctaaaaat	ttaggcgttg	atacggataa	cttatatatt	tctcaaccgg	240
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ttatagtaat	agatagcgtg	gcagctytaa	caccaaaaag	cgaaatagaa	ggcgatatgg	360
gagatcagca	cgtagggctg	caagcaagac	tcatgagtcg	agcacttaga	aaattaaccg	420
gagttgtcca	taaaatggga	actacagttg	tattttataa	ccaaattcgt	atgaaaatcg	480
gcgctatggg	ctatggcact	cctgaaaacta	ctactggcgg	aaatgcgctt	aaattttacg	540
cttcagttgg	acttgacgta	cgtaaaaatg	ctacttttaa	acagagcgat	gagccaatcg	600
gaaaccgcgt	aaaagtaaaa	gtagtaaaaa	acaaagtcgc	tcctcctttt	agacaagccg	660
aatttgatat	catgttttga	gaaggtatca	gcaaagaagg	agagataata	gattacggcg	720

taaaacttga tattatcgat aaaagcggcg cttggtttag ctatgataat tcaaaattag 780
gtcaaggcag agaaaaactca aaagcgtttt taaa 814

<210> 1868

<211> 824

<212> DNA

<213> *Campylobacter jejuni* subsp. *jejuni* ATCC 33560

<400> 1868

```
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cacattatcg cagaatgccca aaaagcagggt ggggtttgtg cttttatcga tgcagaacat 180
gcacttgatg tgaaatatgc taaaaattta ggtgtaaata cagatgattt gtatgtttct 240
caacctgatt ttggagagca agccttagaa attgtagaaa ctatagcwag aagtgggtgca 300
gtagatctta twgtagttaga tagcgttgca gcwcttacc caaaagcaga aattgaaggc 360
gatatgggag atcarcatgt aggacttcaa gcaagactta tgtctcaagc tctaagaaaa 420
cttacaggta tagttcataa aatgaatacc acagtaattt tcatcaacca aattcgtatg 480
aaaatcggtg ctatgggtta tgggtactcct gaaaccacaa cagggtgaaa tgcattaaaa 540
ttttatgctt ctgtgcgttt agatgttaga aaagtagcaa ccttaaamca aaacgwagam 600
cctataggaa accgcgttaa agtaaaagta gttaaaaata aagttgctcc tccattcagm 660
caagctgaat ttgatgtgat gtttggagag ggtttaagcc gtgaagggtga attgatcgat 720
tatggtgtaa aacttgatat cgtagataaa agtgggtgct ggttttctta taaagataaa 780
aaacttggac aaggttagaga aaattcaaaa gctttcttaa aaga 824
```

<210> 1869

<211> 388

<212> DNA

<213> *Enterococcus avium* ATCC 14025

<400> 1869

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agaaagtctt ggtaaaacaa cggttgcact gcatgcgatt gcagaagttc aaaaacatgg 60
ggggacggca gcctttattg atgccgagca cgcgttggac cctcaatcgc cacaacgtct 120
agggtgaaac attgatgaat tgctgctatc acaaccagat actggggaac aaggcttaga 180
aattgcagat gctttagttt caagtggcgc agtcgatatt atcgttattg actcgggtggc 240
cgcgctagtc ccccgtagtg aaatcgatgg cgagatgggt gatgcgcacg ttggtctgca 300
ggctcgtttg atgtcacaag cattgcgcaa gctgtcaggc tctatcaaca aaacaaagac 360
tatcgccgtc ttattaacc aaattcgt 388
```

<210> 1870

<211> 388

<212> DNA

<213> *Enterococcus faecium* ATCC 19434

<400> 1870

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tgaaagtcca ggtaaaacaa cagttgcact acacgctatt gcagaagtac aaaaaaatgg 60
cggaacggcc gctttcattg atgctgagca tgcgttagat ccgcaatatg cacaaaaatt 120
agggtgtgaat atcgatgaac tacttctttc acagcctgac acaggagAAC aaggcttaga 180
gatcgctgat gcttttagtat caagtggggc tgtagatata gtagtagtcg attcagttgc 240
tgcttttagtt ccacgagcag aaatcgacgg cgaaatgggt gactcacatg tcgggttaca 300
agcacgtttg atgtctcaag cattgcgtaa actctctggg tcgatcaaca aaacaaaaac 360
aatcgctatt ttcataacc aaatccgt 388
```

<210> 1871

<211> 388

<212> DNA

<213> *Listeria monocytogenes* ATCC 15313

<400> 1871

```
agagagtccc ggtaaaacaa ctgttgcgct tcatgcaatt gcggaagtac aagcacaagg 60
cggaacagca gcatttatcg atgctgagca tgcgttggat ccggcttatg ctaaaaacct 120
agggtgaaat attgatgaat tattactatc tcaaccagat acaggagAAC aagcttttaga 180
gattgctgaa gcttttagtta gaagtgggtc agttgatatg ttagtaattg actccgttgc 240
```

agcacttgta	ccacgtgctg	aaatcgaagg	cgagatgggc	gatgctcatg	ttggattaca	300
agcacgttta	atgtcccaag	cattgcgtaa	actttctggt	gttattaata	aatcaaaaac	360
cattgctatt	ttcatcaacc	aaattcgt				388

<210> 1872

<211> 388

<212> DNA

<213> *Streptococcus mitis* ATCC 49456

<400> 1872

agagtcac	ggttaagaca	cggttgccct	tcatgcagtt	gcgcaagcac	aaaaagaagg	60
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tgggtgcaac	attgacgaat	tgctcttgtc	acaaccagac	tcaggagagc	aaggtcttga	180
gattgcagga	aaattgattg	actcaggagc	cgtggatctt	gtcgtagtcg	actcagttgc	240
ggcccttgct	cctcgtgcgg	aaattgatgg	agatatcggt	gatagccacg	ttgggttgca	300
ggctcgtatg	atgagccagg	ctatgcgtaa	acttggtgct	tctatcaata	aaacccaaac	360
aattgccatc	tttatcaacc	aattgcgt				388

<210> 1873

<211> 430

<212> DNA

<213> *Streptococcus oralis* ATCC 35037

<400> 1873

gaacatgacg	ccgacttttt	cacgcaattg	gttgataaag	atggcaattg	ttttggtttt	60
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gtgactgtct	ccaatatccc	catcaatttc	cgcacgaggt	acaagggccg	caactgagtc	180
gataacgaca	aggtcaactg	cacctgagtc	aatcaatttt	ccagcaattt	caagaccttg	240
ttcacctgag	tctgggtgtg	acaagagcaa	ttcgtcaata	ttcacaccaa	gggctgcagc	300
ataggctggg	tcaagagcat	gttccgcac	gataaaggct	gcaataccac	cttctttctg	360
tgcttgcgca	acagcgtgaa	gggcaaccgt	tgtcttacca	gatgattctg	gcgcrtaac	420
ttcgtatgata						430

<210> 1874

<211> 947

<212> DNA

<213> *Aspergillus fumigatus* ATCC 64746

<400> 1874

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taaaacacgt	cgcttacggt	ttcgcgaata	ggcccagac	tcgtgagcat	ttgctgctcg	120
cccgccaggt	tgggtgccag	aagatcggtg	tcttcgtcaa	caaaatcgat	gctattgatg	180
atccggagat	gctggaactg	gtcgaactcg	agatgcgtga	gctgctgaac	agctacggtt	240
tcgaggggtga	agagactccg	atcattttcg	gttccgctct	ctgtgctctc	gaaggacgcc	300
gtgacgacat	cggtaaagac	agaattgagc	agcttatgaa	cgctgtcgac	acctggatcc	360
ccactcctca	gcgtgacctc	gacaaaacct	tcttgatgtc	tgctgaggaa	gtgttctcta	420
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cgcacaccgt	gtccagatct	tccgagagat	tagcgatata	tgctaatgat	tcatcagacg	840
aggctgctga	cctcagcttc	cctgacggcg	accaatctcg	cagagttatg	cctgggtgaca	900
acgtcgagat	gatcctgaag	acccaccacc	ctggtgctgc	tgaggct		947

<210> 1875

<211> 923

<212> DNA

<213> *Aspergillus versicolor* strain WSA-175

<400> 1875

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aggtcggtgt	ccagaagatt	gttgtgttcg	tcaacaaggt	tgatgccgtc	gatgaccctg	180
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agaagcactg	tgacgaatcc	cgtgctggtg	acaactccgg	tcttcttctc	cgtggtatcc	600
gccgtgagga	ggtcaagcgc	ggtatggtta	tgtctgctcc	cgccctctatc	aaggcccaca	660
agaagtccat	ggtctccatg	tacgtcctca	ctgaggcaga	aggtggccgt	cgagtggtct	720
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cgacctctcg	ccttagaaga	agaatacctc	taacttgat	ttagacgagg	cttgcgacct	840
ttctttcccc	gatggcgaca	tgagccgccg	tgtcatgcct	ggtgacaacg	tggaaatgat	900
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<210> 1876

<211> 807

<212> DNA

<213> Basidiobolus ranarum ATCC 24670

<400> 1876

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tgatcctgaa	atgttgaggt	tggtcgagat	ggaaatgcgt	gatttgcttt	cccaatacgg	180
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ccccaccctt	gcccgtgatt	tggacaaacc	tttctctcatg	cccgttgaag	atgtgttctc	360
catctctggc	cgttggtactg	ttgccactgg	acgtgttgag	cgtggtatgg	tcaccaaggg	420
tactgaagtt	gaaatcgteg	gtatgggcga	gcacttcaag	accaccttga	ccggtattga	480
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catccttaag	cacccccctg	gtacccccga	tgctgatgag	aagatggtca	tgcccggaga	780
caacgttcaa	ctcgagtgcg	agctctt				807

<210> 1877

<211> 806

<212> DNA

<213> Campylobacter gracilis ATCC 33236

<400> 1877

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ggatctttta	gagttagttg	aagaggaagt	tagagatctt	ttaaaagagt	ataaatcc	180
tggcgacgaa	accccaatca	ttaaggggtc	tgctcttaag	gctcttgagg	aagctaaggc	240
cggacaagac	ggcgaatggt	ctgcaaagat	tatggagctt	atggacgcgg	ttgatagcta	300
tattccaact	cctgttcgcg	atactgataa	agatttctct	cttccgatcg	aagatatttt	360
ctcgatttcc	ggtcgcggtg	ccgttgtaac	cggtagaatc	gaaaaaggta	tcgttaaagt	420
tggtgatact	atcgagatcg	taggtattaa	acctaactcag	actactaccg	tcactggcgt	480
tgagatgttt	agaaaagaga	tggatcaagg	tgaagccggc	gataatgtag	gtgttttatt	540
gcgcggtact	aagaaagagg	aagtagagcg	cggtatgggt	ttatgcaaac	caaaatcgat	600
cactcctcat	actaaatttg	agggcgaggt	ttatatccta	actaaagaag	aaggcggacg	660
ccatactcca	ttctttaata	attatagacc	gcagttttac	gttcgtacga	cagatgttac	720
cggttcgtat	actcttctcg	aaggaaccga	gatgggttatg	ccgggcgaca	acgttaaaat	780
caccgttgag	ctaategctc	cgatcg				806

<210> 1878

<211> 806

<212> DNA

<213> Campylobacter jejuni subsp. jejuni ATCC 33292

```

<400> 1878
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ccaactcgtg atactgaaaa agacttcttg atgccaattg aagatgtttt ctcaatttca 360
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atcgaaatcg ttggtattaa agatactcaa acaacaactg taacagggtg tgaaatgttc 480
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aaaaaagaag aagttatccg tggtatgggt ctgtctaaac caaaatcaat tactccacac 600
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ttctttaaca actatagacc acagttttat gtaagaacaa ctgatgttac aggttcgatt 720
aaattagctg atgggtgtga aatggttatg ccagggtgaaa atgtgagaat tactgtaagc 780
ttgatcgctc cagtagcact tgaaga                                     806

```

<210> 1879

<211> 896

<212> DNA

<213> *Coccidioides immitis* strain WSA-222

```

<400> 1879
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ctcaaaactcg agagcattta cttctcgccc gtcagatcgg tatccaaaaa atcgctcgtc 120
tcgtgaacaa gggtgatgcc atcgaggaca aagagatggt ggagcttggt gaattggaga 180
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tcttgcaggc cgtcgacacc tggattccca ctctcagcg tgagactgac aagcccttct 360
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tggagcgtgg tatcctcaag aaggactccg aagttgaaat tgtcggcggg tcgcccagac 480
caatcaaaac caaggttacc gatatcgaaa cttttaagaa gtcttgcgac gagtctcgcg 540
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aattctatag acgaagcggc tcagctcagc tggcccggag aagatcaaga caagatggct 840
atgccaggag acaatatcga aatgatttgc accaccttgc acccagttgc cgccga 896

```

<210> 1880

<211> 798

<212> DNA

<213> *Erwinia amylovora* ATCC 14976

```

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gaagagctgc tggagctggg tgaaatggaa gtmcgtgacc tgctgtcaca gtacgacttc 180
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gagcgtgcga ttgacaaacc gttcctgctg ccaattgaag acgtgttctc catctctggc 360
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gaaatcgttg gtatcaaaga taccgtgaaa tcaacctgta ccggcgttga gatgttccgt 480
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ttcaaaggct accgtccaca gttctacttc cgtactactg acgtgaccgg gactatcgaa 720
ctgccagaag cggttgagat ggtgatgcca ggcgacaaca ttcagatggt tgtgaccctg 780
atccacccga tcgccatg                                     798

```

<210> 1881

<211> 810

<212> DNA

<213> *Salmonella choleraesuis* subsp. *choleraesuis* ATCC 14028

```
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gacgaagagc tgctggaact ggttgaaatg gaagtccgyg aactgctgtc tcagtacgac 180
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gcagagtggg aagcgaaaat catcgaactg gctggcttcc tggattctta catyccggaa 300
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gttgaaatcg ttggtatcaa agagactcag aagtctacct gtactggcgt tgaaatgttc 480
cgcaaactgc tggacgaagg ccgtgccggt gagaacgtag gtgttctgct gcgtggtatc 540
aaacgtgaag aaatcgaacg tggtcaggta ctggctaagc cgggcacccat caagccgcac 600
accaagttcg aatctgaagt gtacattctg tccaaagatg aaggcggccg tcatactccg 660
ttcttcaaag gctaccgtcc gcagttctac ttccgtacta ctgacgtgac tggcaccatc 720
gaactgccgg aaggcgtaga gatggtaatg ccgggcgaca acatcaaaat ggttggttacc 780
ctgatccacc cgatcgcgat ggacgacggt                                     810
```

```
<210> 1882
<211> 888
<212> DNA
<213> Staphylococcus cohnii strain BM10711
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```
<400> 1882
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ctagatcaga gtggtaggat aaaagaattc gaagtcccta cccctgatgc taaagtgatg 180
tgtttaattg tatcttctact tggagacata tggtttacag agaattggtgc aaataaaaatc 240
ggaaaagctct caaaaaaagg tggctttaca gaatatccat tgccacagcc ggattctggt 300
ccttacggaa taacggaagg tctaaatggc gatatatggt ttaccaat gaattggagat 360
cgtataggaa agttgacagc tgatgggact atttatgaat atgatttgcc aaataaggga 420
tcttatcctg cttttattac tttagggttcg gataacgcac tttggttcac ggagaaccaa 480
aataattcta ttggaaggat tacaatatca gggaaattag aagaatatcc tctaccaaca 540
aatgcagcgg ctccagtggg tatcactagt ggtaacgat gtgcactctg gtttgtcgaa 600
attatgggca acaaaatagg tcgaatcact acaactgggt agattagcga atatgatatt 660
ccaaactcaa acgcacgtcc acacgtata accgcgggga aaaatagcga aatatggttt 720
actgaatggg gggcaaatca aatcggcaga attacaaacg acaaaacaat tcaagaatat 780
caacttcaaa cagaaaatgc ggaacctcat ggtattacct ttggaaaaga tggatccgta 840
tggtttgcat taaaatgtaa aattgggaag ctgaatttga acgaatga                                     888
```

```
<210> 1883
<211> 23
<212> DNA
<213> Artificial Sequence
```

```
<220>
<223> Description of Artificial Sequence:
Oligonucleotide
```

```
<400> 1883
agccgcttga gcaaattaaa cta                                     23
```

```
<210> 1884
<211> 23
<212> DNA
<213> Artificial Sequence
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```
<220>
<223> Description of Artificial Sequence:
Oligonucleotide
```

```
<400> 1884
gtatcccgca gataaatcac cac                                     23
```

<210> 1885
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1885
agcgaaaaac accttgccga c 21

<210> 1886
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1886
gacgcccgcg ccaccact 18

<210> 1887
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1887
gacgcccgcg acaccacta 19
<210> 1888
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1888
gacgcccgca acaccacta 19

<210> 1889
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1889
gttcgcaact gcagctgctg 20

<210> 1890
<211> 19
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1890

ttcgcaacgg cagctgctg

19

<210> 1891

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (14)..(14)

<223> n represents a modified base

<220>

<221> modified_base

<222> (14)..(14)

<223> i

<400> 1891

ccggagctgc cgancggg

18

<210> 1892

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1892

cggagctgcc aarcgggg

18

<210> 1893

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1893

ggagctggcg arcggggt

18

<210> 1894

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:

Oligonucleotide

<400> 1894
gaccggagct agcgarcg 18

<210> 1895
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1895
cggagctagc aarcggggt 19

<210> 1896
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1896
gaaacggaac tgaatgaggc g 21

<210> 1897
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1897
cattaccatg ggcgataaca g 21

<210> 1898
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1898
ccattaccat gagcgataac ag 22

<210> 1899
<211> 861
<212> DNA
<213> Klebsiella pneumoniae strain 15571

<400> 1899
atgcgttata ttgcgctgtg tattatctcc ctggttagcca ccctgccgct ggcggtacac 60
gccagcccgc agccgcttga gcaaattaaa ctaagcgaaa gccagctgtc gggccgcgta 120

ggcatgatag	aaatggatct	ggccagcggc	cgcacgctga	ccgcctggcg	cgccgatgaa	180
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gatgccggtg	acgaacagct	ggagcgaaa	atccactatc	gccagcagga	tctggtggac	300
tactcgccgg	tcagcgaaaa	acaccttgcc	gacggcatga	cggtcggcga	actctgcgcc	360
gccgccatta	ccatgagcga	taacagcgcc	gccaatctgc	tactggccac	cgtcggcgcc	420
cccgcaggat	tgactgcctt	tttgcgccag	atcggcgaca	acgtcaccgc	ccttgaccgc	480
tgggaaacgg	aactgaatga	ggcgcttccc	ggcgacgccc	gcgacaccac	taccccgccc	540
agcatggccg	cgaccctgcg	caagctgctg	accagccagc	gtctgagcgc	ccgttcgcaa	600
cggcagctgc	tgacgtggat	ggtggacgat	cggtcgcgcg	gaccgttgat	ccgtccgtg	660
ctgccggcgg	gctggtttat	cgccgataag	accggagctg	gcgagcgggg	tgcgcgcggg	720
attgtcgccc	tgcttggccc	gaataacaaa	gcagagcgca	ttgtggtgat	ttatctgcgg	780
gataccccgg	cgagcatggc	cgagcgaaat	cagcaaateg	ccgggatcgg	cgcggcgctg	840
atcgagcact	ggcaacgcta	a				861

<210> 1900

<211> 780

<212> DNA

<213> *Klebsiella pneumoniae* strain SLK-47

<400> 1900

ctgttagcca	ccctgccgct	ggcggtacac	gccagcccgc	agccgcttga	gcaaattaaa	60
ctaagcgaaa	gccagctgtc	gggcccgcgta	ggcatgatag	aaatggatct	ggccagcggc	120
cgcacgctga	ccgcctggcg	cgccgatgaa	cgctttccca	tgatgagcac	ctttaaagta	180
gtgctctgcg	gcgcagtgtc	ggcgcggtg	gatgccggtg	acgaacagct	ggagcgaaa	240
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gacggcatga	cggtcggcga	actctgcgcc	gccgccatta	ccatgagcga	taacagcgcc	360
gccaatctgc	tactggccac	cgtcggcgcc	cccgcaggat	tgactgcctt	tttgcgccag	420
atcggcgaca	acgtcaccgc	ccttgaccgc	tgggaaacgg	aactgaatga	ggcgcttccc	480
ggcgacgccc	gcgccaccac	taccccgccc	agcatggccg	cgaccctgcg	caagctgctg	540
accagccagc	gtctgagcgc	ccgttcgcaa	cggcagctgc	tgacgtggat	ggtggacgat	600
cggtcgcgcg	gaccgttgat	ccgtccgtg	ctgccggcgg	gctggtttat	cgccgataag	660
accggagctg	gcgagcgggg	tgcgcgcggg	attgtcgccc	tgcttggccc	gaataacaaa	720
gcagagcgca	ttgtggtgat	ttatctgcgg	gataccccgg	cgagcatggc	cgagcgaaat	780

<210> 1901

<211> 861

<212> DNA

<213> *Escherichia coli*

<400> 1901

atgcgttata	ttcgctgtg	tattatctcc	ctgttagcca	ccctgccgct	ggcggtacac	60
gccagcccgc	agccgcttga	gcaaattaaa	ctaagcgaaa	gccagctgtc	gggcccgcgta	120
ggcatgatag	aaatggatct	ggccagcggc	cgcacgctga	ccgcctggcg	cgccgatgaa	180
cgctttccca	tgatgagcac	ctttaaagta	gtgctctgcg	gcgcagtgtc	ggcgcggtg	240
gatgccggtg	acgaacagct	ggagcgaaa	atccactatc	gccagcagga	tctggtggac	300
tactcgccgg	tcagcgaaaa	acaccttgcc	gacggcatga	cggtcggcga	actctgcgcc	360
gccgccatta	ccatgagcga	taacagcgcc	gccaatctgc	tactggccac	cgtcggcgcc	420
cccgcaggat	tgactgcctt	tttgcgccag	atcggcgaca	acgtcaccgc	ccttgaccgc	480
tgggaaacgg	aactgaatga	ggcgcttccc	ggcgacgccc	gcaacaccac	taccccgccc	540
agcatggccg	cgaccctgcg	caagctgctg	accagccagc	gtctgagcgc	ccgttcgcaa	600
cggcagctgc	tgacgtggat	ggtggacgat	cggtcgcgcg	gaccgttgat	ccgtccgtg	660
ctgccggcgg	gctggtttat	cgccgataag	accggagctg	gcgagcgggg	tgcgcgcggg	720
attgtcgccc	tgcttggccc	gaataacaaa	gcagagcgca	ttgtggtgat	ttatctgcgg	780
gataccccgg	cgagcatggc	cgagcgaaat	cagcaaateg	ccgggatcgg	cgcggcgctg	840
atcgagcact	ggcaacgcta	a				861

<210> 1902

<211> 861

<212> DNA

<213> *Klebsiella pneumoniae* strain 803

<400> 1902

atgcgttata	ttcgctgtg	tattatctcc	ctgttagcca	ccctgccgct	ggcggtacac	60
------------	-----------	------------	------------	------------	------------	----

```

gccagcccg  agccgcttga  gcaaattaaa  caaagcgaaa  gccagctgtc  gggccgcgta  120
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cgctttccca  tgatgagcac  ctttaaagta  gtgctctgcg  gcgcagtgtc  ggcgcgggtg  240
gatgccgggtg  acgaacagct  ggagcgaaaag  atccactatc  gccagcagga  tctggtggac  300
tactcgccgg  tcagcgaaaa  acaccttgcc  gacggcatga  cggtcggcga  actctgcgcc  360
gccgccatta  ccatgagcga  taacagcgcc  gccaatctgc  tgctggccac  cgtcggcgcc  420
cccgcaggat  tgactgcctt  tttgcgccag  atcggcgaca  acgtcaccgc  ccttgaccgc  480
tgggaaacgg  aactgaatga  ggcgcttccc  ggcgacgccc  gcgacaccac  taccgccgcc  540
agcatggccg  cgaccctgcg  caagctgctg  accagccagc  gtctgagcgc  ccgttcgcaa  600
cggcagctgc  tgcagtggat  ggtggacgat  cgggtcgccg  gaccgttgat  ccgctccgtg  660
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attgtcgccc  tgcttgcccc  gaataacaaa  gcagagcgca  ttgtggtgat  ttatctgcgg  780
gatacgccgg  cgagcatggc  cgagcgaaat  cagcaaatcg  ccgggatcgg  cgcggcgctg  840
atcgagcact  ggcaacgcta  a

```

<210> 1903
 <211> 896
 <212> DNA
 <213> *Klebsiella pneumoniae* ATCC 700603

```

<400> 1903
atgcgttatt  ttgcgctgtg  tattatctcc  ctgttagcca  ccctgccgct  ggcggtacac  60
gccagcccg  agccgcttga  gcaaattaaa  ctaagcgaaa  gccagctgtc  gggcagcgta  120
ggcatgatag  aaatggatct  ggccagcggc  cgcacgctga  ccgcctggcg  cgccgatgaa  180
cgctttccca  tgatgagcac  ctttaaagta  gtgctctgcg  gcgcagtgtc  ggcgcgggtg  240
gatgccgggtg  acgaacagct  ggagcgaaaag  atccactatc  gccagcagga  tctggtggac  300
tactcgccgg  tcagcgaaaa  acaccttgcc  gacggcatga  cggtcggcga  actctgtgcc  360
gccgccatta  ccatgagcga  taacagcgcc  gccaatctgc  tgctggccac  cgtcggcgcc  420
cccgcaggat  tgactgcctt  tttgcgccag  atcggcgaca  acgtcaccgc  ccttgaccgc  480
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cggcagctgc  tgcagtggat  ggtggacgat  cgggtcgccg  gaccgttgat  ccgctccgtg  660
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attgtcgccc  tgcttgcccc  gaataacaaa  gcagagcgga  ttgtggtgat  ttatctgcgg  780
gatacgccgg  cgagcatggc  cgagcgaaat  cagcaaatcg  ccgggatcgg  cgcggcgctg  840
atcgagcact  ggcaacgcta  acccggcggt  ggccgcgcgc  gttatccggc  tcgtag      896

```

<210> 1904
 <211> 861
 <212> DNA
 <213> *Escherichia coli* strain JC2926

```

<400> 1904
atgcgttata  ttgcgctgtg  tattatctcc  ctgttagcca  ccctgccgct  ggcggtacac  60
gccagcccg  agccgcttga  gcaaattaaa  ctaagcgaaa  gccagctgtc  gggccgcgta  120
ggcatgatag  aaatggatct  ggccagcggc  cgcacgctga  ccgcctggcg  cgccgatgaa  180
cgctttccca  tgatgagcac  ctttaaagta  gtgctctgcg  gcgcagtgtc  ggcgcgggtg  240
gatgccgggtg  acgaacagct  ggagcgaaaag  atccactatc  gccagcagga  tctggtggac  300
tactcgccgg  tcagcgaaaa  acaccttgcc  gacggcatga  cggtcggcga  actctgcgcc  360
gccgccatta  ccatgagcga  taacagcgcc  gccaatctgc  tactggccac  cgtcggcgcc  420
cccgcaggat  tgactgcctt  tttgcgccag  atcggcgaca  acgtcaccgc  ccttgaccgc  480
tgggaaacgg  aactgaatga  ggcgcttccc  ggcgacgccc  gcgacaccac  taccgccgcc  540
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cggcagctgc  tgcagtggat  ggtggacgat  cgggtcgccg  gaccgttgat  ccgctccgtg  660
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atcgagcact  ggcaacgcta  a

```

<210> 1905
 <211> 861
 <212> DNA
 <213> *Pseudomonas aeruginosa*

```

<400> 1905
atgcgttata ttgcctgtg tattatctcc ctggttagcca ccctgccgct ggcggtacac 60
gccagcccgcc agccgcttga gcaaattaaa ctaagcgaaa gccagctgtc gggccgcgta 120
ggcatgatag aaatggatct ggccagcggc cgcacgctga ccgcctggcg cgccgatgaa 180
cgctttccca tgatgagcac ctttaaagta gtgctctgcy gcgcagtgtc ggcgcgggtg 240
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tactcgccgg tcagcgaaaa acaccttgcc gacggcatga cggtcggcga actctgcgcc 360
gccgccatta ccatgagcga taacagcgcc gccaatctac tactggccac cgtcggcgcc 420
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agcatggccg cgaccctgcy caagctgctg accagccagc gtctgagcgc ccgttcgcaa 600
cggcagctgc tgcagtggat ggtggacgat cgggtcgccg gaccgttgat ccgctccgtg 660
ctgccggcgg gctggtttat cgccgataag accggagcta gcaaacgggg tgcgcgcggg 720
attgtcgccc tgcttgcccc gaataacaaa gcagagcgca ttgtggtgat ttatctgcgg 780
gatacgccgg cgagcatggc cgagcgaaa cagcaaatcg ccgggatcgg cgcggcgctg 840
atcgagcact ggcaacgcta a 861

```

```

<210> 1906
<211> 20
<212> DNA
<213> Artificial Sequence

```

```

<220>
<223> Description of Artificial Sequence:
      Oligonucleotide

```

```

<400> 1906
ccttattccc ttttttgccg                                     20

```

```

<210> 1907
<211> 22
<212> DNA
<213> Artificial Sequence

```

```

<220>
<223> Description of Artificial Sequence:
      Oligonucleotide

```

```

<400> 1907
cacctatctc agcgatctgt ct                                     22

```

```

<210> 1908
<211> 23
<212> DNA
<213> Artificial Sequence

```

```

<220>
<223> Description of Artificial Sequence:
      Oligonucleotide

```

```

<400> 1908
aacagcggta agatccttga gag                                     23

```

```

<210> 1909
<211> 22
<212> DNA
<213> Artificial Sequence

```

```

<220>
<223> Description of Artificial Sequence:
      Oligonucleotide

```

<400> 1909
atgacttggt taagtactca cc 22

<210> 1910
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1910
atgacttggt tgagtactca cc 22

<210> 1911
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1911
ccataaccat gggtgataac ac 22

<210> 1912
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1912
ccataaccat gagtgataac ac 22

<210> 1913
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1913
cgccttgatc attgggaacc 20

<210> 1914
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1914

cgcccttgatc gttgggaacc 20

<210> 1915
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1915
cgcccttgata gttgggaacc 20

<210> 1916
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1916
cgtgggtctt gcggtatcat 20

<210> 1917
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1917
cgtgggtctg gcggtatcat 20

<210> 1918
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1918
gtgggtctca cggtatcatt g 21

<210> 1919
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1919
cgtgggtctc tcggtatcat t 21

<210> 1920
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
<221> misc_feature
<222> (6)..(6)
<223> n represents any nucleotide

<400> 1920
cgtggnctctc gcggtatcat 20

<210> 1921
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1921
cgtgggtcta gcggtatcat t 21

<210> 1922
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1922
gttttccaat gattagcact ttta 24

<210> 1923
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1923
gttttccaat gataagcact ttta 24

<210> 1924
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:

Oligonucleotide

<400> 1924
gtttttccaat gctgagcact ttt 23

<210> 1925
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1925
cgtttttccaa tgatgagcac ttt 23

<210> 1926
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1926
gtttttccaat ggtgagcact ttt 23

<210> 1927
<211> 861
<212> DNA
<213> Neisseria meningitidis strain MC9690-129

<400> 1927
atgagtattc aacatttttcg tgtcgccctt attccctttt ttgcggcatt ttgccttcct 60
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tgcagtgcgt ccataaccat gagtgataac actgctgcca acttacttct gacaacgatc 420
ggaggaccga aggagctaac cgcttttttg cacaacatgg gggatcatgt aactcgcctt 480
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tcccggcaac aattaataga ctggatggag gcggataaag ttgcaggacc acttctgcgc 660
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cgcggtatca ttgcagcact ggggccagat ggtaagccct cccgtatcgt agttatctac 780
acgacgggga gtcaggcaac tatggatgaa cgaaatagac agatcgctga gatagggtgcc 840
tactgatta agcattggta a 861

<210> 1928
<211> 861
<212> DNA
<213> Escherichia coli strain HB251

<400> 1928
atgagtattc aacattttccg tgtcgccctt attccctttt ttgcggcatt ttgccttcct 60
gttttttgctc acccagaaac gctggtgaaa gtaaaagatg ctgaagatca gttgggtgca 120
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cgtggtgacg ccgggcaaga gcaactcggc cgccgcatac actattctca gaatgacttg 300

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gttaagtact caccagtcac agaaaagcat cttacggatg gcatgacagt aagagaatta 360
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cgcggtatca ttgcagcact ggggccagat ggtaagccct cccgtatcgt agttatctac 780
acgacgggga gtcaggcaac tatggatgaa cgaaatagac agatcgctga gatagggtgcc 840
tcaactgatta agcattggta a                                     861

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<210> 1929

<211> 808

<212> DNA

<213> *Klebsiella oxytoca* strain 26W

<400> 1929

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ggaggaccga aggagctaac cgcttttttg cacaacatgg gggatcatgt aacctgcctt 480
gatagttggg aaccggagct gaatgaagcc ataccaaacg acgagcgtga caccacgatg 540
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tcggcccttc cggtggctg gtttattgct gataaatctg gagccggtga gcgtgggtct 720
cgcggtatca ttgcagcact ggggccagat ggtaagccct cccgtatcgt agttatctac 780
acgacgggga gtcaggcaac tatggatg                                     808

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<210> 1930

<211> 861

<212> DNA

<213> *Escherichia coli*

<400> 1930

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atgagtattc aacatttttcg tgtcgccctt attccctttt ttgcggcatt ttgccttct 60
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cgagtgggtt acatcgaact ggatctcaac agcggtaaga tccttgagag ttttcgcccc 180
gaagaacgtt ttccaatgat gagcactttt aaagtctctg tatgtggcgc ggtattatcc 240
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gttgagtact caccagtcac agaaaagcat cttacggatg gcatgacagt aagagaatta 360
tgcagtgtct ccataaccat gagtgtataac actgctgcca acttacttct gacaacgatc 420
ggaggaccga aggagctaac cgcttttttg cacaacatgg gggatcatgt aactcgctt 480
gatcgttggg aaccggagct gaatgaagcc ataccaaacg acgagcgtga caccacgatg 540
cctgcagcaa tggcaacaac gttgcgcaaa ctattaactg gcgaactact tactctagct 600
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ggcggtatca ttgcagcact ggggccagat ggtaagccct cccgtatcgt agttatctac 780
acgacgggga gtcaggcaac tatggatgaa cgaaatagac agatcgctga gatagggtgcc 840
tcaactgatta agcattggta a                                     861

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<210> 1931

<211> 861

<212> DNA

<213> *Escherichia coli* strain BM2728

<400> 1931

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cgagtgggtt acatcgaact ggatctcaac agcggtaaga tccttgagag ttttcgcccc 180

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gaagaacggt	ttccaatgat	gagcactttt	aaagtctctg	tatgtggcgc	ggtattatcc	240
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 <212> DNA
 <213> Escherichia coli

<400> 1932						
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cgagtgggtt	acatcgaaat	ggatctcaac	agcggtaaga	tccttgagag	ttttcgcccc	180
gaagaacggt	ttccaatgct	gagcactttt	aaagtctctg	tatgtgggtg	ggtattatcc	240
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gttgagtact	caccagtcac	agaaaagcat	cttacggatg	gcatgacagt	aagagaatta	360
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tcactgatta	agcattggta	a				861

<210> 1933
 <211> 861
 <212> DNA
 <213> Escherichia coli

<400> 1933						
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tcactgatta	agcattggta	a				861

<210> 1934
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Oligonucleotide

<400> 1934
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<210> 1935
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1935
agctcggcat acttcgacag g 21

<210> 1936
<211> 15
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1936
taccacccgc acggc 15

<210> 1937
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1937
cggagtcgcc gtcgatg 17

<210> 1938
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1938
ccgcgcacca ttgcttcgta cactgaggag tctccgcgcg g 41

<210> 1939
<211> 46
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:

Oligonucleotide

<400> 1939
cgacccggat ggtagtatcg ataatgatcc gccagcggcc gggtcg 46

<210> 1940
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1940
gtatcgttgg tgacgtaat 19

<210> 1941
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
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<400> 1941
gcaatggtcc gtttaagt 18

<210> 1942
<211> 27
<212> DNA
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<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1942
gactggaaca aagcctataa aaaatca 27

<210> 1943
<211> 16
<212> DNA
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<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1943
gctggtggac ggccag 16

<210> 1944
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1944
tttcgccgcc atgcgttac 19

<210> 1945
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1945
cggcgactac gcggtat 17

<210> 1946
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1946
cggcgacttc gcggtat 17

<210> 1947
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
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<400> 1947
cggtatacgg caccatcgt 19

<210> 1948
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1948
gcggtataca acaccatcg 19

<210> 1949
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1949

cggtatacgc caccatcgt 19

<210> 1950
 <211> 15
 <212> DNA
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<220>
 <223> Description of Artificial Sequence:
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<400> 1950
 ggcgacatcg cctgc 15

<210> 1951
 <211> 17
 <212> DNA
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<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1951
 ggcgacagag cctgcta 17

<210> 1952
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1952
 cctgctatgg agcgatggt 19

<210> 1953
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 1953
 cgcctgctat aaagcgatgg t 21

<210> 1954
 <211> 589
 <212> DNA
 <213> Klebsiella pneumoniae subsp. pneumoniae ATCC 13883

<400> 1954
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 ctttacgcc tgaacgtatt gggcaatgac tggaaacaaag cctataaaaa atcagcccgt 180
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tttgggttcca	tcgacggcga	ctccgcgcgcg	gcgatgcggtt	ataccgaaat	tcgtctggcg	360
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aacggcgcc	cgggatcgc	cgtagggatg	gccaccaaca	taccgccaca	taacctgacg	540
gaagtgatta	acggctgtct	ggcgtatgtt	gacgatgaag	acatcagca		589

<210> 1955

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1955

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<210> 1956

<211> 989

<212> DNA

<213> Candida inconspicua ATCC 16783

<400> 1956

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agaagttgaa	attgtttggtg	gtaaagaagg	tgttattaag	actactgtta	ccggtattga	720
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ggcatacaag	aagttcttat	catcattata	cattttaaca	aaggaagaag	gtggttagaca	900
tactccattt	tctgaaaatt	acagacctca	aatgtacatt	agaacttcca	atgttaattgt	960
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<210> 1957

<211> 991

<212> DNA

<213> Candida utilis ATCC 22023

<400> 1957

ggtaagacca	cccttactgc	cgccatcacc	aagtgccttg	ctgagaaggg	aggtgcctcg	60
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agtgcctttg ctgagaacta cagaccacag atgttcatca gaaccggaga tgtcaccacc 960
atcttgacat ggccagagga gcacgtgac c 991

<210> 1958

<211> 985

<212> DNA

<213> *Candida zeylanoides* ATCC 7351

<400> 1958

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ctcgactgcc	cacgttgagt	acgagaccga	taagagacac	tacgcccacg	ttgattgccc	180
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<210> 1959

<211> 973

<212> DNA

<213> *Candida catenulata* ATCC 10565

<400> 1959

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<210> 1960

<211> 985

<212> DNA

<213> *Candida krusei* ATCC 28870

<400> 1960

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catcaagatt	atattaagaa	tatgattact	ggtgctgcac	aaatggatgg	tgctattatt	240
gttggttctg	ctactgatgg	tcaaagtcca	caaactaagg	aacatttatt	attagcaaga	300
caagttgggtg	ttcaacattt	agttgtcttt	gttaataaat	gtgacaccat	tgatgaccca	360

gaaatggttg	aattagttga	aatggaaatg	agagaactat	tgtctgaata	tggttttgat	420
ggtgataaca	ctccagttat	tatgggttct	gcattgatgg	ctttagaaga	caagagacct	480
gaagttggta	aggaatctat	tttaaagtta	atggaagcyg	ttgacacatg	gattccaacc	540
ccagagagag	atttagaaaa	accatttttg	ttacctattg	atgaagtttt	ctcaatctct	600
ggtagaggta	ctgtcgtttc	tggtactgtc	gaaagaggta	ctttgaagaa	gggtgaagaa	660
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tatcacaagg	aattagacca	agcgcaagca	ggtgatactc	caggtatttt	attaagaggt	780
gtcaagagag	accaaataca	gagaggtcaa	attttagcaa	agccagattc	cgtaaggca	840
tacaagaagt	tcttggcttc	cctttatatc	ttaaccaagg	aagaagggtg	tagacataca	900
ccattctctg	aaaactacag	accacaaatg	tacatcagaa	ctaccaatgt	taacgttact	960
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<210> 1961
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1961
gctcaaggca gatggcattc cc 22

<210> 1962
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1962
ggacaaggcg gttgcgtttg at 22

<210> 1963
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1963
cattcccgtc tcgctcgaca gt 22

<210> 1964
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1964
atctgcctgc ccgtcttgc 19

<210> 1965

<211> 816
<212> DNA
<213> Unknown Organism

<220>
<223> Description of Unknown Organism: Plasmid pGS05

<400> 1965
atgaataaat cgctcatcat tttcggcatc gtcaacataa cctcggacag tttctccgat 60
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ggggcagatg tgatcgacct cggtcgggca tccagcaatc ccgacgccgc gcctgtttcg 180
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gcacccgctg gcgacatcat ggatcacatt gcggcgttct ttgacgcgcg catcgcggcg 480
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gggtggagctg acttcatccg cacacacgag ccgcgccct tgcgcgacgg gctggcggtg 780
ttggcgcgcg tgaaagaaac cgcaagaatt cgtaa 816

<210> 1966
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1966
catgccagtc ttgccaacg 19

<210> 1967
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1967
cagcaataag taatccagcg atg 23

<210> 1968
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1968
ggagagattt caccgcatag 20

<210> 1969
<211> 22
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1969

agccaacat catgctattc ca

22

<210> 1970

<211> 1206

<212> DNA

<213> Unknown Organism

<220>

<223> Description of Unknown Organism: Transposon Tn10

<400> 1970

atgaatagtt	cgacaaagat	cgcattggta	attacgttac	tcgatgccat	ggggattggc	60
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aaccactttg	gcgtattgct	tgcactttat	gcgttaatgc	aggttatctt	tgctccttgg	180
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gcacgcgtgg	attacttatt	gctggctttt	tcaagtgcgc	tttggatgct	gtatttaggc	300
cgtttgcttt	cagggatcac	aggagctact	ggggctgtcg	cggcatcggt	cattgccgat	360
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gaaacaaaaa	atacacgtga	taatacacat	accgaagtag	gggttgagac	gcaatcgaat	600
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caattgatag	gccaaattcc	cgcaacgggt	tggtgtgctat	ttaccgaaaa	tcgttttgga	720
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gtgatgtcta	tccaaacaaa	gagtcatgag	caagggtgct	tacagggatt	attgggtgagc	1020
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ctaccaattt	gggatggctg	gatttggatt	attggtttag	cgttttactg	tattattatc	1140
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<210> 1971

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1971

cygactgygc catcctyadc a

21

<210> 1972

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (3)..(3)
<223> n represents a modified base

<220>
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<222> (13)..(13)
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<221> modified_base
<222> (3)..(3)
<223> i

<220>
<221> modified_base
<222> (13)..(13)
<223> i

<400> 1972
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21

<210> 1973
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
<221> misc_feature
<222> (9)..(9)
<223> n represents a modified base

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<222> (16)..(16)
<223> n represents a modified base

<220>
<221> modified_base
<222> (9)..(9)
<223> i

<220>
<221> modified_base
<222> (16)..(16)
<223> i

<400> 1973
racaccrgny ttggwntcct t

21

<210> 1974
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
<221> misc_feature

<222> (8)..(8)
<223> n represents a modified base

<220>
<221> modified_base
<222> (8)..(8)
<223> i

<400> 1974
acaaggntg grmsaaggag ac 22

<210> 1975
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1975
tgrccrgggt ggttraggac g 21

<210> 1976
<211> 21
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence:
Oligonucleotide

<220>
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<222> (16)..(16)
<223> n represents a modified base

<220>
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<222> (16)..(16)
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<400> 1976
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<210> 1977
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1977
gatggaytcy gtyaartggg a 21

<210> 1978
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
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Oligonucleotide

<220>
<221> misc_feature
<222> (5)..(5)
<223> n represents a modified base

<220>
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<222> (5)..(5)
<223> i

<400> 1978
catcntgyaa tggyaatcty aat

23

<210> 1979
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1979
catcytgyaa tggyaascty aat

23

<210> 1980
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
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<222> (9)..(9)
<223> n represents a modified base

<220>
<221> misc_feature
<222> (12)..(12)
<223> n represents a modified base

<220>
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<222> (14)..(14)
<223> n represents a modified base

<220>
<221> modified_base
<222> (9)..(9)
<223> i

<220>
<221> modified_base
<222> (12)..(12)
<223> i

<220>

<221> modified_base
<222> (14)..(14)
<223> i

<400> 1980
tcratggcnt cnanragrgt yt

22

<210> 1981
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
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<222> (9)..(9)
<223> n represents a modified base

<220>
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<222> (15)..(15)
<223> n represents a modified base

<220>
<221> modified_base
<222> (9)..(9)
<223> i

<220>
<221> modified_base
<222> (15)..(15)
<223> i

<400> 1981
tggacaccns caagnggkcy g

21

<210> 1982
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
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<222> (9)..(9)
<223> n represents a modified base

<220>
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<222> (15)..(15)
<223> n represents a modified base

<220>
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<222> (9)..(9)
<223> i

<220>
<221> modified_base
<222> (15)..(15)
<223> i

<400> 1982
tggacacyns caagnggkcy g 21

<210> 1983
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
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<222> (14)..(14)
<223> n represents a modified base

<220>
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<222> (14)..(14)
<223> i

<400> 1983
cygaytgcg yatnctcatc a 21

<210> 1984
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1984
cygaytgygc yatyctsatc a 21

<210> 1985
<211> 1383
<212> DNA
<213> Cryptococcus neoformans strain M1-106

<400> 1985
atgggtaagg acaagctgca cgtcaacgtc gttgttatcg gtcacgtcga ctccggtaag 60
tcgaccacca ccggctcactt gatctacaag tgcgggtggta tgcacaagcg aaccattgag 120
aagttcgaga aggaggctca agagctcggg aagtcttctt tcaagtacgc ttgggttctt 180
gacaagctta aggccgagcg agagcgaggt atcaccatcg acattgctct ttggaagtgc 240
gagaccacctt agtaccaggt taccgtcatt gacgcccccg gtcaccgaga cttcatcaag 300
aacatgatca ccggtacctc ccaggctgac tgtgccatcc tcatcattgc caccggatatc 360
ggtgagttcg aggctggtat ctccaaggac ggtcagaccc gagagcacgc ctcctcgcgc 420
ttcaccctcg gtgtcaggca gctcattggt gcttgaaca agatggacac ctgcaagtgg 480
tctgaggacc gattcaacga aatcgtcaag gagaccaacg gtttcatcaa gaagggttgg 540
tacaacccca aggetgtccc ctctcgtccc atctctggtt ggcacggtga caacatgttg 600
gaggagacca ccaacatgcc ctggtacaag ggatggacca aggagaccaa gtccgggtgtt 660
tccaagggtg agacccttct cgaggccatc gacgccagta ggccccctac ccgacccacc 720
gacaagcccc tccgtctccc tctccaggac gtctacaaga tcggtggtat cggcacagtc 780
cctgtcggcc gagtcgagac cgggtgtcatc aaggccggta tggtcgtcaa gttcgcctcc 840
accaacgtca cactgaagt caagtccgtt gagatgcacc acgagcagat ccccgagggt 900

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aagcttggtt	cccagaagcc	tctctgtggt	gagacctacg	ccgactaccc	ccctcttggt	1260
cgattcgccg	tccgagacat	gcgacagacc	gttgccgttg	gtgttattaa	gagtggtggag	1320
aagtccgatg	ggaagagcgg	caaggttacc	aaggccgccg	agaaggctgc	taagaagaag	1380
taa						1383

<210> 1986

<211> 1380

<212> DNA

<213> *Cryptococcus neoformans* strain B3501

<400> 1986

atgggtaagg	acaagctgca	cgtcaacgtc	gttggttatcg	gtcacgtcga	ctccggtaag	60
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aagttcgaga	aggaggctca	agagctcggg	aagctcttctt	tcaagtacgc	ttgggttctt	180
gacaagctta	aggccgagcg	agagcgaggt	atcaccatcg	acattgctct	ttggaagttc	240
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aagcttggtg	cccagaagcc	cctctgtggt	gagacctacg	ccgactaccc	ccctcttggt	1260
cgattcgccg	tccgagacat	gcgacagacc	gttgccgttg	gtgttatcaa	gagcgtggac	1320
aagaccgaga	agggtggcaa	ggtcaccaag	gctgctgaga	aggctgccaa	gaagaagtaa	1380

<210> 1987

<211> 1377

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 1987

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aagttcgaaa	aggaagccgc	tgaattaggt	aagggttctt	tcaagtacgc	ttgggttttg	180
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tctggtaaga	agttggaaga	ccatccaaag	ttcttgaagt	ccggtgacgc	tgctttgggtc	1200
aagttcggtc	catctaagcc	aatgtgtgtt	gaagctttca	gtgaataccc	accattaggt	1260
agattcgctg	tcagagacat	gagacaaact	gtcgtgtcgc	gtgttatcaa	gtctgttgac	1320
aagactgaaa	aggccgctaa	ggttaccaag	gctgctcaaa	aggctgctaa	gaaataa	1377

<210> 1988

<211> 1377

<212> DNA

<213> *Saccharomyces cerevisiae*

<400> 1988

atgggtaaaag	agaagtctca	cattaacggt	gtcggttatcg	gtcatgtcga	ttctggtaag	60
tctaccacta	ccggtcattt	gatttacaag	tgtggtggta	ttgacaagag	aaccatcgaa	120
aagttcgaaa	aggaagccgc	tgaattaggt	aagggttctt	tcaagtacgc	ttgggttttg	180
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gaaactccaa	agtaccaagt	taccgttatt	gatgctccag	gtcacagaga	tttcatcaag	300
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<210> 1989

<211> 1377

<212> DNA

<213> *Eremothecium gossypii* ATCC 10895

<400> 1989

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<210> 1990
<211> 1377
<212> DNA
<213> *Eremothecium gossypii*

<400> 1990
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<210> 1991
<211> 1646
<212> DNA
<213> *Aspergillus oryzae* strain KBN616

<400> 1991
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<210> 1992

<211> 1380

<212> DNA

<213> Aureobasidium pullulans strain R106

<400> 1992

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<210> 1993

<211> 1383

<212> DNA

<213> Histoplasma capsulatum strain 186AS

<400> 1993

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<210> 1994
<211> 1383
<212> DNA
<213> *Neurospora crassa*

<400> 1994
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<210> 1995
<211> 1383
<212> DNA
<213> *Podospira anserina*

<400> 1995
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<210> 1996
<211> 1386

<212> DNA

<213> *Podospora curvicolla* strain VLV

<400> 1996

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<210> 1997

<211> 1383

<212> DNA

<213> *Sordaria macrospora* strain 000

<400> 1997

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<210> 1998

<211> 1383

<212> DNA

<213> *Trichoderma reesei* strain QM9414

<400> 1998
atgggtaagg aggacaagac tcacatcaac gtgggtcgta tcggccacgt cgactccggc 60
aagtctacca ccactgggtca cttgatctac cagtgcggtg gtatcgacaa gcgtaccatt 120
gagaagttcg agaaggaagc cgccgaactc ggcaagggtt ccttcaagta cgcgtgggtt 180
cttgacaagc tcaaggccga gcgtgagcgt ggtatcacca tcgacattgc cctctggaag 240
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aagaacatga tcaactggtac ttcccaggcc gactgcgcta tcctcatcat cgctgccggg 360
actggtgagt tcgaggctgg tatctccaag gatggccaga cccgtgagca cgctctgctc 420
gcctacaccc tgggtgtcaa gcagctcatc gtcgccatca acaagatgga cactgccaac 480
tgggccgagg ctcgttacca ggaaatcatc aaggagactt ccaacttcat caagaaggtc 540
ggcttcaacc ccaaggccgt tgccttcgtc cccatctccg gcttcaacg tgacaacatg 600
ctcacccctt ccaccaactg cccctggtac aagggtcggg agaaggagac caaggctggc 660
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gtcaagatga tcccctccaa gcccattgtc gttgaggctt tcaccgacta ccctcccctg 1260
ggtcgtttcg ccgtccgtga catgcgccag accgtcgctg tcggtgtcat caaggccgct 1320
gagaagtcct ctgccgccgc cgccaaggtc accaagtccg ctgccaaggc cgccaagaaa 1380
taa

<210> 1999
<211> 29
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 1999
catgtcaaya ttggtactat tgggtcatgt 29

<210> 2000
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
<221> misc_feature
<222> (9)..(9)
<223> n represents a modified base

<220>
<221> modified_base
<222> (9)..(9)
<223> i

<400> 2000
ccaccytcnc tcamgttgaa rcggt 25

<210> 2001
<211> 23
<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (6)..(6)

<223> n represents a modified base

<220>

<221> misc_feature

<222> (12)..(12)

<223> n represents a modified base

<220>

<221> modified_base

<222> (6)..(6)

<223> i

<220>

<221> modified_base

<222> (12)..(12)

<223> i

<400> 2001

acyacnttra cngcygcyat yac

23

<210> 2002

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (3)..(3)

<223> n represents a modified base

<220>

<221> misc_feature

<222> (15)..(15)

<223> n represents a modified base

<220>

<221> modified_base

<222> (3)..(3)

<223> i

<220>

<221> modified_base

<222> (15)..(15)

<223> i

<400> 2002

ccngargara gagcnmgwgg t

21

<210> 2003

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (9)..(9)

<223> n represents a modified base

<220>

<221> modified_base

<222> (9)..(9)

<223> i

<400> 2003

catytcranr ttgtcacctg g

21

<210> 2004

<211> 1360

<212> DNA

<213> Candida albicans strain SC5314

<400> 2004

gctgccttcg	accgtttctaa	acctcatgtc	aacattggta	ctattggtca	tgttgatcat	60
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ttcttggatt	atggttctat	tgatagagct	ccagaagaaa	gagctagagg	tatcactatt	180
tccactgccc	acgttgaata	cgaaaccaag	aacagacact	atgcccacgt	tgattgtcca	240
ggacacgctg	attatatcaa	aaatatgatt	actggtgccg	ctcaaatgga	tggtgctatc	300
attgttggtg	ctgccactga	tggtcaaatg	cctcaaacca	gagaacattt	gttattggcc	360
agacaagtgt	gtgttcaaga	cttggttgtg	tttgtcaaca	aagtcgatac	tattgatgac	420
cctgaaatgt	tggaattagt	cgaaatggaa	atgagagaat	tggtatccac	ctacggtttt	480
gatggtgaca	acactccagt	tattatggga	tctgctttaa	tggctttgga	agacaagaaa	540
ccagaaattg	gtaaggaagc	tatcttgaaa	ttgttagatg	ctgtcgatga	acacattcca	600
actccatcaa	gagacttgga	acaaccattt	ttgttaccag	ttgaagacgt	gttctccatc	660
tccggtagag	gaactgttgt	cactggtaga	gttgaaagag	gtgttttgaa	gaagggtgaa	720
gaaatcgaaa	ttgttgggtg	ttttgacaaa	ccttacaaga	ctactgttac	cggtattgaa	780
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tctcacaaga	agttcttggc	tcccttgat	atcttgactt	ccgaagaagg	tggtcgttcc	960
actccatttg	gtgaagggtt	caagcctcaa	tgcttcttca	gaactaacga	tgctactacc	1020
acattttcat	tcccagaagg	agaagggtgt	gatcattctc	aatgatcat	gccagggtgac	1080
aacattgaaa	tggttggtga	attgatcaaa	tcttgtccat	tagaagtcaa	ccaacgtttc	1140
aacttgagag	aagggtggtaa	aactgttggt	actggtttga	ttaccagaat	catcgaataa	1200
acagaatgtg	cactgtgaat	aataaaaaga	aaagagggtat	atatagggtga	ctttgtattt	1260
tgtattgaac	aataaaaattc	tgtaaatagt	aagggcctca	gaagttttga	tttgatttat	1320
gccatgtgga	cttgtagaga	tatccttctc	aaacttcttg			1360

<210> 2005

<211> 1342

<212> DNA

<213> Schizosaccharomyces pombe

<400> 2005

aagccgcatg	tcaatattgg	tactattggt	catgttgacc	acggtaaaac	gacgttgacg	60
gctgctatta	ctaaatgcct	ttctgatctt	gggtcaagcta	gttttatgga	ttatagtcaa	120
attgacaagg	cccccgagga	aaaggcacgt	gggtattacca	tttcatctgc	ccatggtgaa	180
tacgaaactg	ctaactcgta	ctatgcccct	gtggattgtc	ctgggtcacgc	cgattacatt	240
aagaatatga	ttactgggtg	tgctacaatg	gatggcgcta	tcattgttgt	ttctgtacc	300
gatgggtcaa	tgccatcaaa	tcgtgaacat	ttgcttctgg	ctcgtcaagt	cggtgtaaa	360
caaattgttg	tatacatcaa	taaagtcgat	atgggtcgagc	ctgatatgat	cgagcttgtc	420
gaaatggaaa	tgcggtgagct	actctccgaa	tacggatttg	atgggtgacaa	tactccaatt	480
gttagcggca	gtgctttatg	tgctttagag	gggtcgtagc	ctgagattgg	tctcaatagt	540


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attactaaat  tgatggaagc  tgttgatagt  tatattactc  ttcctgaaag  aaaaacggat  600
gtccctttct  tgatggccat  cgaggacgtt  ttttcaattt  caggtcgcgg  aactgtagtc  660
actggccgtg  tcgagcgcg  tactttaaag  aagggtgctg  aaatcgaaat  cgtcggttat  720
ggtagccatt  taaagactac  cgttactgga  attgaaatgt  tcaaaaagca  gcttgatgcc  780
gccgttgccg  gtgacaattg  tggcctttta  cttcgttcta  tcaagcgaga  gcaattaaaa  840
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ttctatattt  tgacaaaaga  ggaaggaggt  cgtcgtaccg  gtttcggtga  caagtatcgt  960
ccccaactgt  acagtcgtac  ttccgacgtt  actgtcgaac  ttaccacccc  tgatcctaac  1020
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cccattgtca  tcgaaaaagg  acaacgcttc  acagttcggt  aggggtggaag  cactgtaggc  1140
acagctttgg  ttactgaact  tttggattag  tgcatttatg  aacttattgg  ctttaaaaaat  1200
tttgcattgt  gaataccaat  attatgtccc  ttctcagaat  tctataacta  cagtgtcatt  1260
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attgtagcc  aaagttataa  aa  1342

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<210> 2006
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 2006
 tggagccggt gagcgtgg 18

<210> 2007
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 2007
 tggagccagt gagcgtgg 18

<210> 2008
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 2008
 tctggagccg atgagcgtg 19

<210> 2009
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 2009
 ctggagccag taagcgtgg 19

<210> 2010
<211> 861
<212> DNA
<213> *Klebsiella pneumoniae* strain KMK107

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<400> 2010
atgagtattc aacatttttcg tgtcgccctt attccctttt ttgcggcatt ttgccttcct 60
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cgagtgggtt acatcgaaact ggatctcaac agcggtaaga tccttgagag ttttcgcccc 180
gaagaacggt ttccaatgat gagcactttt aaagttctgc tatgtggtgc ggtattatcc 240
cgtgttgacg ccgggcaaga gcaactcggg cgccgcatac actattctca gaatgacttg 300
gttaagtact caccagtcac agaaaagcat cttacggatg gcatgacagt aagagaatta 360
tgcagtgtcg ccataacccat gagtataaac actgctgcca acttacttct gacaacgac 420
ggaggaccga aggagctaac cgcttttttg cacaacatgg gggatcatgt aactcgccct 480
gatcggtggg aaccggagct gaatgaagc ataccaaagc acgagcgtga caccacgacg 540
cctgcagcaa tggcaacaac gttgcgcaaa ctattaactg gcgaactact tactctagct 600
tcccggcaac aattaataga ctggatggag gcggataaag ttgcaggacc acttctgcgc 660
tcggcccttc cggctggctg gtttattgct gataaatctg gagccagtga gcgtgggtct 720
cgcggtatca ttgcagcact ggggccagat ggtaagccct cccgtatcgt agttatctac 780
acgacgggga gtcaggcaac tatggatgaa cgaaatagac agatcgctga gatagggtgcc 840
tcactgatta agcattggta a                                     861
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<210> 2011
<211> 861
<212> DNA
<213> *Klebsiella pneumoniae* strain CLSiS L-491

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<400> 2011
atgagtattc aacattttccg tgtcgccctt attccctttt ttgcggcatt ttgctttcct 60
gtttttgctc acccagaaac gctggtgaaa gtaaaagatg ctgaagatca gttgggtgca 120
cgagtgggtt acatcgagct ggatctcaac agcggtaaga tccttgagag ttttcgcccc 180
gaagaacggt ttccaatgat gagcactttt aaagttctgc tatgtggtgc ggtattatcc 240
cgtgttgacg ccgggcaaga gcaactcggg cgccgcatac actattctca gaatgacttg 300
gttgagtact caccagtcac agaaaagcat cttacggatg gcatgacagt aagagaatta 360
tgcagtgtcg ccataacccat gagtataaac actgcggcca acttacttct gacaacgac 420
ggaggaccga aggagctaac cgcttttttg cacaacatgg gggatcatgt aaccgcctt 480
gatcggtggg aaccggagct gaatgaagc ataccaaagc acgagcgtga caccacgatg 540
cctgcagcaa tggcaacaac gttgcgcaaa ctattaactg gcgaactact tactctagct 600
tcccggcaac aattaataga ctggatggag gcggataaag ttgcaggacc acttctgcgc 660
tcggcccttc cggctggctg gtttattgct gataaatctg gagccagtaa gcgtggatct 720
cgcggtatca ttgcagcact ggggccagat ggtaagccct cccgtatcgt agttatctac 780
atgacgggga gtcaggcaac tatggatgaa cgaaatagac agatcgctga gatagggtgcc 840
tcactgatta agcattggta a                                     861
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<210> 2012
<211> 49
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<220>
<221> misc_feature
<222> (29)..(29)
<223> n represents a modified base

<220>
<221> modified_base
<222> (29)..(29)
<223> n = guanidyl-MR-HEG

<400> 2012
ccgcggatta ttaaaccgcc cttccgcgna tgtcagaggg atagatcca 49

<210> 2013
<211> 353
<212> DNA
<213> Kluyvera ascorbata ATCC 33433

<400> 2013
agcttaagaa ctcttatctg gattacgcga tgtcgggtcat tggtggccgt gcgctgccgg 60
atgtccgaga tggcctgaag ccggtacacc gtcgcgtact ttacgccatg aacgtattgg 120
gcaatgactg gaacaaagcc tacaaaaaat cagcccgtgt cgtgggtgac gtgatcggta 180
aatatcaccc gcatgggtgat actgccgtct atgacactat cgtccgtatg gcacagccat 240
tctcactgcg atacatgctg gtagatggtc aaggtaactt cggttctgtc gatggcgact 300
ccgcgcgacg gatgcgttat acggaaatcc gtatgtcgaa aatcgcccat gag 353

<210> 2014
<211> 355
<212> DNA
<213> Kluyvera georgiana ATCC 51603

<400> 2014
agctcctatc tggattatgc gatgtcggtc attggtggcc gtgcgctgcc agatgtccga 60
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gcaatgcggt atacggaaat ccgtctggcg aaaattgccc atgaactgat ggccg 355

<210> 2015
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2015
ccaagaagct caaaaacatc tg 22

<210> 2016
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2016
tadcctgtcc awacagccat 20

<210> 2017
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2017
actttgaata aggtcgggtct ag 22

<210> 2018
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2018
acactaaaca aggttggttt ag 22

<210> 2019
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2019
acactaaaca aggtcgggtct ag 22

<210> 2020
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2020
gtagctccag atgaaatggt tg 22

<210> 2021
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2021
gtagctccag acgaaatggt tg 22

<210> 2022
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2022

gtagctccag atgaaacggtt tg 22

<210> 2023
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2023
gtaactccag atgaaatggtt tg 22

<210> 2024
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2024
agtgaaaaga tggctgctgc 20

<210> 2025
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2025
agtgagaaaa tggctgctgc 20

<210> 2026
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2026
tccaagcatg cattatgcaa acg 23

<210> 2027
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2027
tcggtctaga tagagctaaa acg 23

<210> 2028
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2028
tatgctcttc aacaatcacg 20

<210> 2029
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2029
agccgttgag actttgaata ag 22

<210> 2030
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2030
cttaatgggc ttggtatcg 19

<210> 2031
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2031
cgtgactggg gttctgctat ga 22

<210> 2032
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2032
cgtgactggg gatcatcaat ga 22

<210> 2033
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2033
cgtgactggg gttctgccat ga 22

<210> 2034
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2034
atcaagaaca ctggctatgt ag 22

<210> 2035
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2035
atcaagaaca ctggctacgt ag 22

<210> 2036
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2036
atcaagaaca ctggttacgt ag 22

<210> 2037
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2037
atcaaaaata ctggttatgt ag 22

<210> 2038

<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2038
atcaagaata ctggctacgt ag 22

<210> 2039
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2039
atcaaaaaca ctggctatgt ag 22

<210> 2040
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2040
tgtgacccca gacaaaccc 19

<210> 2041
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2041
gttgagcggc agcactatct 20

<210> 2042
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2042
cacggggatt tctctattta 20

<210> 2043
<211> 20

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2043
cacggggatt actctatttta 20

<210> 2044
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2044
accgtaagtc ggccaagtca 20

<210> 2045
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2045
gttcttttctc cgtatcgtc 19

<210> 2046
<211> 20
<212> DNA
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<211> 2160

<212> DNA

<213> Streptococcus pneumoniae strain R6

<400> 2048

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<212> DNA

<213> Streptococcus pneumoniae strains 63509; M11

<400> 2051

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<211> 1195

<212> DNA

<213> Streptococcus pneumoniae strain #22/HA5

<400> 2052

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<211> 930

<212> DNA

<213> Streptococcus pneumoniae strain 17619

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<210> 2054

<211> 306

<212> DNA

<213> Streptococcus pneumoniae strain R6

<400> 2054

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<211> 2472

<212> DNA

<213> Streptococcus pneumoniae strain 7785

<400> 2055

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<210> 2056

<211> 1212

<212> DNA

<213> Streptococcus pneumoniae strain StrR-16

<400> 2056

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<210> 2057

<211> 1242

<212> DNA

<213> Streptococcus pneumoniae strain StrR-17

<400> 2057

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<210> 2058

<211> 1225

<212> DNA

<213> Streptococcus pneumoniae strain StrR-18

<400> 2058

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<210> 2059

<211> 554

<212> DNA

<213> Streptococcus pneumoniae strain StrR-38

<400> 2059

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acaagcaggt aagacaggta cttctaacta tactgacgaa gaaattgaaa agtatatcaa 480
gaatgctggg tacgtagctc cagatgaaat gtttgttggg tatacccgca aatatgcaat 540
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<210> 2060

<211> 1249

<212> DNA

<213> Streptococcus pneumoniae strain StrR-57

<400> 2060

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atgttttctt cggaattaac caagcagtag aaacaaaccg cgactgggga tcaactatga 240
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<210> 2061

<211> 579

<212> DNA

<213> Streptococcus pneumoniae strain StrR-60

<400> 2061

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<210> 2062

<211> 1216

<212> DNA

<213> Streptococcus pneumoniae strain StrR-63

<400> 2062

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<210> 2063

<211> 810
<212> DNA
<213> Streptococcus pneumoniae ATCC 700673

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<210> 2064
<211> 782
<212> DNA
<213> Streptococcus pneumoniae ATCC 700678

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<210> 2065
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2065
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<210> 2066
<211> 22
<212> DNA
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<220>
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<400> 2066

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<210> 2067
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
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<400> 2067 18
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<210> 2068
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<220>
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<400> 2068 19
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<210> 2069
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<210> 2070
<211> 18
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<220>
<223> Description of Artificial Sequence:
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<210> 2071
<211> 18
<212> DNA
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<220>
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<400> 2071 18
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<210> 2072
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 <213> Mycobacterium tuberculosis strain Rv

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<223> Description of Artificial Sequence:
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<210> 2074
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2074
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<210> 2075
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2075
ccgagccagg ttctgaagtc tctgcattat taggtgctcg g 41

<210> 2076
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2076
ccgagcygay aacattttca gattcaccca rgcgctcgg 39

<210> 2077
<211> 33
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2077
ccgagcaacc gatccagctc cagctacgct cgg 33

<210> 2078
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2078
ccgagccttg gtcttcggcc aaatgaacgc tcgg 34

<210> 2079
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2079
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<210> 2080
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<212> DNA
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<220>
<223> Description of Artificial Sequence:
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<400> 2080
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<210> 2081
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2081
gratyrtyaa agttggtgag gaag 24

<210> 2082
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<212> DNA
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<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2082
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<210> 2083

<211> 44
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<220>
<223> Description of Artificial Sequence:
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<222> (27)..(27)
<223> n represents a modified base

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<222> (27)..(27)
<223> i

<400> 2083
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44

<210> 2084
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<212> DNA
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<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2084
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<210> 2085
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2085
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<210> 2086
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<220>
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Oligonucleotide

<400> 2086
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23

<210> 2087
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<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2087
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<210> 2088
<211> 22
<212> DNA
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<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2088
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<210> 2089
<211> 20
<212> DNA
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<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2089
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<210> 2090
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<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2090
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<210> 2091
<211> 39
<212> DNA
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<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2091
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<210> 2092
<211> 38
<212> DNA
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<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2092
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<210> 2093
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<212> DNA
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<220>
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<210> 2094
<211> 24
<212> DNA
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<220>
<223> Description of Artificial Sequence:
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<400> 2094
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<210> 2095
<211> 18
<212> DNA
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<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2095
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<210> 2096
<211> 34
<212> DNA
<213> Artificial Sequence

<220>
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<210> 2097
<211> 1185
<212> DNA
<213> Mycoplasma pneumoniae ATCC 29342

<400> 2097
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<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 2098
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<210> 2099
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 2099
 ccaggactga acgggatacg aa 22

<210> 2100
 <211> 18
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 2100
 gcgagacgat aggttgctc 18

<210> 2101
 <211> 2609
 <212> DNA
 <213> Mycobacterium tuberculosis strain H37Rv

<400> 2101

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<210> 2102
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2102
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20

<210> 2103
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:

Oligonucleotide

<400> 2103
cgaaccagcg gaaatagttg gac

23

<210> 2104
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

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<221> misc_feature
<222> (12)..(12)
<223> n represents a modified base

<220>
<221> modified_base
<222> (12)..(12)
<223> i

<400> 2104
ctgggcatgg cncgagtc

18

<210> 2105
<211> 3297
<212> DNA
<213> Mycobacterium tuberculosis strain H37Rv

<400> 2105
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
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<220>
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<210> 2133
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<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
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<212> DNA
<213> Pseudomonas aeruginosa strain PAO-1

<400> 2138
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<212> DNA
<213> Campylobacter jejuni NCTC 11168

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<210> 2140

<211> 2157

<212> DNA

<213> *Streptococcus pneumoniae* strain 670

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<211> 18
<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2141
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<210> 2142
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<212> DNA
<213> Artificial Sequence

<220>
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Oligonucleotide

<400> 2142
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<210> 2143
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
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Oligonucleotide

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<210> 2144
<211> 3075
<212> DNA
<213> Staphylococcus aureus strain J2870

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aaaatgtctc gagtagaaga agtaatcgat gtttggtttg atagcggctc tatgccgttt 1560
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gaaattacaa aaagtagaaa tacattagat aattgggctc tttctcgctt aaacacctta 2040
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gaatcaataa aaaccaataa aaatatgttt aaagaaaatt tcgtgattaa aaatatacac 3000
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<210> 2145
 <211> 22
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 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

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 gcaagatgtg gcgtgttacg gt 22

<210> 2146
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

<400> 2146
 ggggcgaaga agttgtccat att 23

<210> 2147
 <211> 660

<212> DNA
<213> Escherichia coli

<400> 2147
atggagaaaa aaatcactgg atataccacc gttgatatat cccaatggca tcgtaaagaa 60
cattttgagg catttcagtc agttgctcaa tgtacctata accagaccgt tcagctggat 120
attacggcct ttttaaagac cgtaaagaaa aataagcaca agttttatcc ggcctttatt 180
cacattcttg cccgcctgat gaatgctcat ccggaattcc gtatggcaat gaaagacggg 240
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gacaagggtg tgatgccgct ggcgattcag gttcatcatg ccgtctgtga tggcttccat 600
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<210> 2148
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<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2148
cagattaaat gcggattcag cc 22

<210> 2149
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2149
atcaggtaaa tcatcagcgg ata 23

<210> 2150
<211> 642
<212> DNA
<213> Escherichia coli strain K12

<400> 2150
atgaatttta cccggattga cctgaatacc tggaatcgca gggaacactt tgccctttat 60
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accgcactgg cggagacagg ttataagttt tatccgctga tgatttacct gatctcccgg 180
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tcttgccgtt attttcgga tctcagttag tttatggcag gttataatgc ggtaacggca 360
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atatcatcat taccgtgggt gagttttgac ggatttaacc tgaacatcac cggaaatgat 480
gattattttg ccccggtttt tacgatggca aagtttcagc aggaagggtga ccgcgtatta 540
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attaatacac ttcagctgat gtgtgataac atactgaaat aa 642

<210> 2151
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2151
atatttcagc attaccttgg gtt 23

<210> 2152
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2152
tacacaactc ttgtagccga tta 23

<210> 2153
<211> 642
<212> DNA
<213> Shigella flexneri

<400> 2153
atgaactata caaaatttga tgtaaaaaat tggggttcgcc gtgagcattt tgagttttat 60
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gattcagtcg acccacaatt caccgtattc catcaagaaa cagagacatt ttcagcactg 300
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attaatcggc tacaagagtt gtgtaacagt aaattaaaaat aa 642

<210> 2154
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2154
cgccattcag agtttaggac 20

<210> 2155
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2155
ttccataccg ttgcgtatca ctt 23

<210> 2156
<211> 624
<212> DNA
<213> Clostridium perfringens strain CP590

<400> 2156
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tttgcgaagt taccttgtac atacagcatg accgttaaag tggatatcac acaaataaag 120
gaaaaggga tgaaactata tcctgcaatg ctttattata ttgcaatgat tgtaaaccgc 180
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ataccgtggt caaccttcga tggctttaat ctgaatttgc agaaaggata tgattatttg 480
attcctattt ttactatggg gaaatattat aaagaagata acaaaattat acttcctttg 540
gcaattcaag ttcacacgc agtatgtgac ggatttcaca tttgccgttt tgtaaacgaa 600
ttgcaggaat tgataaatag ttaa 624

<210> 2157
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2157
ccacagaaat tgatattagt gttttat 27

<210> 2158
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2158
tcgctattgt aaccagttct a 21

<210> 2159
<211> 651
<212> DNA
<213> Staphylococcus aureus

<400> 2159
atgaacttta ataaaattga tttagacaat tggaagagaa aagagatatt taatcattat 60
ttgaaccaac aaacgacttt tagtataacc acagaaattg atattagtgt tttataccga 120
aacataaaaac aagaaggata taaattttac cctgcattta ttttcttagt gacaaggggtg 180
ataaaactcaa atacagcttt tagaactggt tacaatagcg acggagagtt aggttattgg 240
gataagttag agccacttta tacaattttt gatggtgtat ctaaaacatt ctctggtatt 300
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ctttctatta ttccatggac ttcatttact gggtttaact taaatatcaa taataatagt 480
aattaccttc taccattat tacagcagga aaattcatta ataaaggtaa ttcaatatat 540
ttaccgctat ctttacaggt acatcattct gtttgtgatg gttatcatgc aggattgttt 600
atgaactcta ttcaggaatt gtcagatagg cctaagtact ggctttttata a 651

<210> 2160
<211> 23

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2160
ttttgaacac tattttaacc agc 23

<210> 2161
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2161
gatttaactt atcccaataa cct 23

<210> 2162
<211> 648
<212> DNA
<213> Staphylococcus aureus

<400> 2162
atgacttttta atattatcaa attagaaaat tgggatagaa aagaatattt tgaacactat 60
tttaaccagc aaactacgta tagcattact aaagaaattg atattacttt gtttaaagat 120
atgataaaaa agaaaggata tgaaatttat ccttctttga tttatgcaat tatggaagtt 180
gtaaataaaa ataaagtgtt tagaacagga attaatagtg agaataaatt aggttattgg 240
gataagttaa atcctttgta tacagttttt aataagcaaa ctgaaaaatt tactaacatt 300
tggactgaat ctgataacaa cttcacttct ttttataata attataaaaa tgacttgctt 360
gaatataaag ataaagaaga aatgtttcct aaaaaaccga tacctgaaaa caccataccg 420
atthcaatga ttccttggat tgatttttagt tcattttaatt taaacattgg taacaatagc 480
aactttttat tgcctattat tacgataggt aaattttata gtgagaataa taaaatttat 540
ataccagttg ctttgcagct tcatcatgct gtatgtgatg gttaccatgc ttcattattt 600
atgaatgaat ttcaagatat aattcataag gtagatgatt ggatttag 648

<210> 2163
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2163
accttcatcc taccgatgtg gggt 24

<210> 2164
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2164

caacgacacc agcactgccca ttg

23

<210> 2165
<211> 1215
<212> DNA
<213> Salmonella typhimurium strain H3380

<400> 2165
atgaccacca cacgccccgc gtgggcctat acgctgccgg cagcactgct gctgatggct 60
ccttttcgaca tcctcgcttc actggcgcgat gatatttata tccctgtcgt tccagcgatg 120
cccggcatcc tgaacacgac gcccgctatg atccaactca cgttgagcct ctatatgggtg 180
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gcggtgacgc tgttaaacgg cgatacagc tggcccgatg tttgttacgc cacggcaatg 1140
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<210> 2166
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2166
gacaaacat tcctgctg 18

<210> 2167
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2167
cagcagctgg gcggcggt 18

<210> 2168
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2168
catcaaagtt ggtgaagaag ttg 23

<210> 2169
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2169
cccgtttgcg aaaggtgg 18

<210> 2170
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2170
acgtgacgtt gacaaacca 19

<210> 2171
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2171
tcgttgatt aactgaagaa 20

<210> 2172
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2172
gtggtgaaat gttccgtaaa 20

<210> 2173
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2173

aagaaaaaat cttcgaactg gcta

24

<210> 2174

<211> 15

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2174

tctacacggc cggtg

15

<210> 2175

<211> 16

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2175

ccgccataacc ccgttt

16

<210> 2176

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2176

cggcattacc atttccacac cttt

24

<210> 2177

<211> 49

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2177

ggcacggaca aaccattcct gctgcctatc gaagacgtgt tcccgtgcc

49

<210> 2178

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2178

ggcacgacaa accattcctg ctgcctatcg aacgtgcc

38

<210> 2179
<211> 39
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2179
ggcagctcta cttccgtacc actgacgtaa ccggctgcc

39

<210> 2180
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<212> DNA
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<220>
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Oligonucleotide

<400> 2180
ttcgccggcg tgggc

15

<210> 2181
<211> 15
<212> DNA
<213> Artificial Sequence

<220>
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Oligonucleotide

<400> 2181
agcgccacgc gcagg

15

<210> 2182
<211> 42
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2182
gcgcgccaac gacttctacc acgaaatgga agagtgcgc gc

42

<210> 2183
<211> 817
<212> DNA
<213> *Alcaligenes faecalis* subsp. *faecalis* ATCC 8750

<400> 2183
tatcttgggtt wgctcggccg ctgacggccc aatgcctcag actcgcgagc acatcctgct 60
gagccgtcag gttggcgttc cttacatcat cgtgttcctg aacaaggccg acatgggttg 120
tgacgaagag ctgatcgaac tggttgaaat ggaagttcgc gagctgttgt ccaagtacga 180
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caccatcgag	ctgccagaag	acaaggaaat	ggttctgcca	ggcgacaaca	tttcgatgaa	780
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<210> 2184

<211> 1652

<212> DNA

<213> *Campylobacter coli* ATCC 43479

<400> 2184

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tccaagaata	gtattttgtaa	ataaaatgga	tagaatcggt	gcaaatttct	acaatgtaga	180
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agaagaatat	cgcacaaaaa	tgatagaagc	agtttctgaa	acttcagatg	agttgatgga	420
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<210> 2185

<211> 820

<212> DNA

<213> *Succinivibrio dextrinosolvens* ATCC 19716

<400> 2185

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cagttcccag	gcgacgacac	tccaatcatc	cgtgggttcag	cactaggtgc	attaaacggc	240
gaagagaagt	ggaaagaggc	aatctatcag	ttagcagaca	ctctagattc	atacattcca	300
gagccaaagc	gtgatatcga	tgatccattc	ctattaccaa	tcgaagatat	cttctcaatc	360
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gaagttgaaa	tcgttggtat	tcgtccaacc	accaagacca	ctgtaactgg	cgttgaaatg	480
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cacaccaagt	tcactgggtca	ggtttacgtg	ctaagcaagg	atgaaggtgg	tcgtcacact	660
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<210> 2186
<211> 44
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
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<400> 2186
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<210> 2187
<211> 1612
<212> DNA
<213> Campylobacter jejuni subsp. jejuni ATCC 33292

<400> 2187
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aagatcaaat tcgcaaccgt ttaaaagcta atccagttcc acttcaaatt ccaatcggtg 180
ctgaggataa ttttaaaggc gtaatcgatc ttgtaactat gaaagcttta gtttgggaag 240
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cagaagaata tcgcacaaaa atgatagaag cagtttctga aacttcagat gagttgatgg 360
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tgaaaactcc tgaagattac atgggtgatg ttattggaga tcttaacaaa cgccgtgggtc 1560
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<210> 2188
<211> 1667
<212> DNA
<213> Campylobacter jejuni subsp. jejuni ATCC 33560

<400> 2188
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ccaagaatag tatttgtaaa taaaatggat agaatcgggtg caaatttcta caatgtagaa 180
gatcaaatc gcaaccgttt aaaagctaatt ccagttccac ttcaaattcc aattgggtgct 240
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gatactaagc caacggatta tgtagaaaaa gaaattccag ctgaacttaa agaaaaggca 360
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caacctttgc ttgatgctgt tgtggcttat ttaccagctc ctgatgaagt ggcaaatatc 600
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gcaggacttg catttaaaat catgacagat ccattttagt gacaacttac tttcgtgcgt 720

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gaaactcctg	aagattatat	gggtgatgtt	attggagatc	ttaacaaacg	ccgtgggtcaa	1620
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<210> 2189

<211> 1255

<212> DNA

<213> Leishmania guyanensis ATCC 50126

<400> 2189

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<210> 2190

<211> 1248

<212> DNA

<213> Trypanosoma brucei subsp. brucei strain EATRO795

<400> 2190

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catcttgacg	cccacacggg	ccgctgtatc	gcgatgcaaa	cgacggatct	cctcaaactg	120
aaggcaaagg	tcgtttcgac	aggtggcaac	atttccgttc	ctgtcggccg	ggaaacacta	180
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<210> 2191

<211> 440

<212> DNA

<213> *Aspergillus nidulans* strain WSA-176

<400> 2191

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gggtaacgat	ctgtaccacg	aaatgcagga	gactgggtgc	attcagctcg	acggcgaatc	180
caaggtgtct	cttgtgttcg	gtcagatgaa	cgagccccc	ggtgctcgtg	ccgtgtcgc	240
ccttactggg	ctgaccatcg	ccgaataactt	ccgtgacgag	gagggtcagg	acgtgctgct	300
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tcgtatcccc	tctgccgtcg	gttaccagcc	cactctggcc	gtcgacatgg	gtgggtatgca	420
ggaacgtatt	accaccacca					440

<210> 2192

<211> 1262

<212> DNA

<213> *Leishmania panamensis* ATCC 50158

<400> 2192

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<210> 2193

<211> 912

<212> DNA

<213> *Aspergillus nidulans* strain WSA-176

<400> 2193

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agaagatcgt	tgtcttcgtc	aacaagggtg	acgctgtcga	tgacctgag	atgttgagc	180
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<210> 2194

<211> 887

<212> DNA

<213> Aureobasidium pullulans strain WSA-234

<400> 2194

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tattgactgg	cccgaaggca	ccgaggacgc	tgactccaag	atggtcatgc	ccggtgacaa	840
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<210> 2195

<211> 984

<212> DNA

<213> Emmonsia parva ATCC 10784

<400> 2195

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<210> 2196

<211> 806

<212> DNA

<213> Exserohilum rostratum strain WSA-215

<400> 2196

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ggcgagaact	acaggccaca	aatgttcatc	cgtactgctg	acgagtcatg	cgcgctgcac	720
tgccagaag	gtacccaga	tgctcacgac	aagcttgta	tgcttggtga	taacgttgag	780
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<210> 2197

<211> 799

<212> DNA

<213> *Fusarium moniliforme* strain WSA-213

<400> 2197

accggtgaac	acttgctcct	cgctcgtcag	gttggtgttc	agcgaattgt	cgtctttgtc	60
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gagcttctta	acacctatgg	cttcgaaggc	gacgacactc	ccgtcatcat	gggctcggct	180
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gctgcgctcg	acgagtggat	cccaaccccc	gagcgtgacc	ttgacaagcc	cttccttatg	300
tccgtcgagg	atgtcttctc	cattgctggc	cgtggtaccg	tcgtgtctgg	ccgtgtggag	360
cgtggtgttc	tgaagcgtga	ccaggagatc	gagcttggtg	gaaaggggtca	ggaggttatc	420
aagaccaagg	ttaccgacat	cgagaccttc	aagaagtctt	gtgagcagtc	ccaggctggg	480
gacaactctg	gtctcctcat	ccgaggtggt	cgccgtgagg	atgtccgccc	tggtatgggc	540
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<210> 2198

<211> 819

<212> DNA

<213> *Fusarium solani* ATCC 32793

<400> 2198

ctctgacggg	cagatgcccc	agacccgtga	gcacttgctg	cttgcccgtc	aggtcggtgt	60
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gctcgtcgag	atggagatgc	gtgagctcct	caacacctac	ggcttcgagg	gtgacgagac	180
ccctgtcatc	atgggctctg	ctctcatgtc	cctccagaac	cagcgccccg	agatcggtag	240
ccagaagatt	gacgagctcc	ttgccgccgt	tgacgagtgg	atccctaccc	ccgagcgtga	300
ccttgacaag	cccttctctc	tgtccgttga	ggatgtcttc	tccattgccg	gccgtgggtac	360
cgctgtctct	ggccgtgtcg	agcgtgggtg	cctgaagcgc	gaccaggaga	ttgagctcgt	420
cggcaagggt	aacgagggtca	tcaagaccaa	ggtcaccgac	attgagacct	tcaagaagtc	480
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ggatgtccgc	cgtggtatgg	tcgtctgcgc	ccccggcact	gtcaagtccc	acactcagtt	600
cctttcttcc	ctctacgtcc	tcaccaagga	ggagggtggc	cgacacactg	gcttccagga	660
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cgagggtacc	gaggacgcca	gcagcaagat	ggtcatgcc	ggtgacaaca	ccgagatggg	780
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<210> 2199

<211> 1025

<212> DNA

<213> *Histoplasma capsulatum* strain WSA-377

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<400> 2199
tgctgacggc caaatgtaag acgccgcgag ggagtgtgta aggttttatg ctttttaggc 60
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cacgtgagca tttgtctcct gcccgacagg tcggtgtcca aaagatcgtc gttttcgtga 180
acaaagtcga cgcccttgag gacaaggaga tgttgagact tgtcgagtta gaaatgagag 240
agctcttaaa cacctacgga tttgagggtg aagagacacc catcatcttt gggtctgccc 300
tttgcgccat ggaaggccgt gagcctgagt tgggagaaaa gaaaattgat gaattgctgg 360
aggctgttga tacttgatc ccaacaccac aacgtgatac cgaaaaacct ttcttgatgt 420
ccgttgagga agtattctct atctccggtc gtggaaccgt tgcctccggt cgtgttgagc 480
gcggtgtcct caagaaggat tcagaagtcg agctaattgg gggcggtcct acccccatca 540
ggacgaaggt aactgatatc gaaactttca agaaatcctg tgacgagtct agagctgggg 600
acaactccgg tcttttattg cgtggtatca agcgtgaaga tatccgccgt ggtatggtag 660
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ccgaagctga ggggtggtcgc cgaaccggat tcggccagaa ctatcgctct caaatgttca 780
tccgcacagc tgggtatgtca aaatggggcc ctttttcata atcctttctt ttttcccttt 840
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cccccttgat tatagacgaa gccgcccatc tcagcttccc tagtggagca gatgaaagca 960
aactcgttat gcctggtgac aacgtcgaga tgatcctcca gacacaccgc cccgtggctg 1020
ctgag                                     1025

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<210> 2200
<211> 667
<212> DNA
<213> Kocuria kristinae ATCC 27570

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<400> 2200
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ccgacatggt cgaggacgag gagctgctgg acctcgctga gatggaggtc cgcgagctgc 120
tgtctctcca ggagttcgac ggcgacaacg ccccgctcat ccgctgaagg gcgctgaagg 180
cgctggaggg cgactgagaag tgggtcaagt ccatcgagga gctcatggag gccgtggacg 240
agtacatccc ggaccccgct cgcgacaagg acaagccgtt cctgatgccc atcgaggacg 300
tcttcaccat caccgggccc ggcaccgtgg tgaccggtcg cgccgagcgc gggaccctgg 360
ccctgaactc cgaggtcgag atcgctcgga tccgcccgat ccagaagacc acggtcaccg 420
ggatcgagat gttccacaag cagctcgacg aggcctgggc cggcgagaac tgcggtctgc 480
tgctgcgcgg cctgaagcgc gacgacgtcg agcgcggcca ggtcgtggtg aagccggggt 540
ccatcacccc gcacaccaac ttcgaggcga acgtctacat cctgtccaag gacgagggtg 600
ggcgtcacia cccgttctac tcgaactacc gtccgcagtt ctacttccgg accaccgacg 660
tcaccgg                                     667

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<210> 2201
<211> 778
<212> DNA
<213> Vibrio mimicus ATCC 33653

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<400> 2201
tgctgcaaca gatggtccaa tgccacaaac tcgtgagcac atcctgctgg gtcgccaaagt 60
aggatttcct tacatcatcg tattcatgaa caaatgtgac atggttgacg atgaagagct 120
tctagagctg gttgagatgg aagttcgtga gcttctgtct gactacgatt tcccagggtga 180
tgacctgcca gtaatccaag gttcagcact aggcgcgcta aacggcgaag cacagtggga 240
agcgaagatt gttgaactag cagaagcact agattcatac attccagagc cagagcgtgc 300
agtagacatg gcattcctga tgccaatcga agacgtattc tcaatccaag gtcgtggtag 360
agtagtaact ggccgtatcg agcgcggcat cctgaaagtg ggtgacgaag ttgcgatcgt 420
tggtatcaaa gacacagtaa aaactacctg tacagggtgta gaaatgttcc gtaagctgct 480
tgacgaaggt cgtgcagggt agaacggttg tgcactgcta cgtggtagta agcgtgaaga 540
agtagagcgt ggtcaagtag tggcgaagcc aggttcaatc accccacaca ctaagttcga 600
atcagaagta tacgtactgt caaaagacga aggtggccgt catactccat tcttcaaaag 660
ttaccgtcca cagttctact tccgtacaac tgacgtaaca ggcagcatcg agcttccaga 720
aggcgtagaa atggtaatgc caggcgacaa catcaagatg gttgtagacc tgattgca 778

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<210> 2202
<211> 412
<212> DNA
<213> Citrobacter freundii ATCC 8090

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<400> 2202
aacgctgacc ctgcagggtta ttgccgctgc gcagcgtgaa ggtaaaacct gtgcattttat 60
cgatgcagaa cacgcactgg acccggtcta tgcccgttaag cttggcggttg atatcgataa 120
cctgctgtgt tctcagccgg ataccggtga acaagcgtg gaaatctgtg atgcaactggc 180
gcgctccggt gcggttgacg ttatcgttgt cgactccgtt gccgcattga cgccgaaggc 240
agaaatcgaa ggcgagattg gcgactctca catgggcctt gcggcgcgta tgatgagcca 300
ggcgatgcgt aagctggccg gtaacctgaa gcagtccaac acgctgctga ttttcatcaa 360
ccagatccgt atgaagattg gcgttatgtt cggtaacccg gaaaccacca cc 412

<210> 2203
<211> 337
<212> DNA
<213> Clostridium botulinum strain 20:1.2

<400> 2203
tttagatcca tcttatgcta gaaatttagg tgttgatata gataacctaa tagttttctca 60
accagataca ggagaacagg ctttagagat aacagaagct ttagtaagat caggagcagt 120
agatgttata gttgtagact ctgtagcagc tttagttcct agggcagaaa tagaaggaga 180
aatgggagac tcacatgtag gtcttcaagc aagacttatg tctcaagccc taagaaaatt 240
agcaggatct ataaataaat ctaagtgtgt agctatatatt ataaaccaat taagagaaaa 300
ggttggtata atgtttggaa atccagaaac aactcct 337

<210> 2204
<211> 379
<212> DNA
<213> Francisella tularensis ATCC 29684

<400> 2204
aaagcaaggc ggtactgcag catttggtga tgetgagcat gcactagatc caaaatatgc 60
aaagctttta ggtgttgatg ttgataatct gatcgtgtca cagccggata cgggtgagca 120
agcttttagag attgctgata tggttggtacg ttctggagga gttgatattg tagtaattga 180
ctctgttget gcacttacgc caaaggcaga gattgagggg gacatgggag actcgacat 240
gggcttacaa gcaagattaa tgtcacaaag actaagaaaa ctaacggcaa atatcaagcg 300
ctcaaatact ctagtgtatg tcattaacca aattcgtatg aagatcgggg ttatgttttg 360
taaccctgaa actacaact 379

<210> 2205
<211> 337
<212> DNA
<213> Peptostreptococcus anaerobius ATCC 27337

<400> 2205
acttgaccca gtatatgcaa gggctcttgg agtggatata gacaacctag tcatatctca 60
gccagataca ggagaacagg ccctagatat agcagagtcc cttataagat caggagctgt 120
agatatacta gtaatagact cagtagctgc ctagtagcct aaggcagaaa tagaagggtga 180
catgggagat tctcacgtag gtctacagcg tagacttatg tcacaggcac ttagaaaaatt 240
gactggatct ataaagaagt caaactgtgt tgttatatatt atcaaccagt tgagagaaaa 300
agtagggggt atgttcggta atccagagac aacaaca 337

<210> 2206
<211> 337
<212> DNA
<213> Peptostreptococcus asaccharolyticus strain LSPQ 2639

<400> 2206
tcttgatgct ggatatgcaa aaaaccttgg agtagatgta gaaaatttaa ttattttctca 60
acctgatata ggtgagcaag ccttagaaat aactgaagct cttgtaagat ctaacgctgt 120
tgatttaatt attatagact cagttgccgc acttgtagca aaagcagaaa tcgatgggtga 180
catgggagct gcacaaatag gtcttcaagc aagacttatg tctcaagctc ttagaaaaatt 240
aactggggca atcaacaagt caaaatgtac cgttgtattt attaaccaac ttagagaaaa 300
agttggtatc atgtttggta acccagaaac tacaaca 337

<210> 2207
 <211> 408
 <212> DNA
 <213> *Providencia stuartii* ATCC 33672

<400> 2207
 ctcacgttgc aagttattgc agcagcacaa cgtagcggaa aaacctgtgc atttatcgac 60
 gctgaacatg cgctagatcc aatctatgcg aaaaaactgg gtgttgatat cgataacctt 120
 ctatgttctc aacctgatac tggtagagca gcattagaga tttgtgatgc actgacgcgt 180
 tcaggcgctg ttgatgtcat tatcgttgac tccgtggccg cattaacacc aaaagctgaa 240
 attgaagggtg aaatcgggtga ctcacacatg ggcttagcgg ctcgatatgat gagccaagcg 300
 atgcgtaaat tagcgggtaa cttaaagaac tcgaatacac ttttaattctt cattaaccaa 360
 atccgtatga agattggcgt tatgtttggt aaccagaaa ccactaca 408

<210> 2208
 <211> 403
 <212> DNA
 <213> *Salmonella choleraesuis* subsp. *choleraesuis* ATCC 9150

<400> 2208
 gctgcagggtg attgccgctg cgcagcgtga aggtaaaacc tgtgcgttta tcgatgcgga 60
 acacgcgctt gaccctgttt acgcacgcaa gctgggcgtc gatatcgata acctgctttg 120
 ttctcagccg gataccggcg agcaggcgct ggaaatctgt gacgcgctgg cgcgttcagg 180
 cgcggtggac gtcattgttg tcgactccgt agcggcgcta acgccgaaag cggaatcga 240
 aggcgaaatt ggcgactctc acatgggcct cgcggcgctg atgatgagcc aggcgatgcg 300
 taagctggcg gggaacctaa aacagtccaa cacgctgttg attttcatca accagatccg 360
 tatgaagatt ggcgtgatgt tcggtaaccc ggaaaccacc acc 403

<210> 2209
 <211> 412
 <212> DNA
 <213> *Salmonella choleraesuis* subsp. *choleraesuis* ATCC 14028

<400> 2209
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 gcgttcaggc gcggtggacg tcattgtggt cgactccgta gcggcgctaa cgccgaaagc 240
 ggaaatcgaa ggcgaaatcg gcgactctca catgggcctc gcggcgcgta tgatgagcca 300
 ggcgatgcgt aagctggcgg ggaacctgaa acagtccaac acgctgttga ttttcatcaa 360
 ccagatccgt atgaagattg gcgtgatgtt cggtaacccg gaaaccacca cc 412

<210> 2210
 <211> 337
 <212> DNA
 <213> *Staphylococcus saprophyticus* ATCC 15305

<400> 2210
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 acctgatcat ggggaacaag gtttagaaat tgccgaagca tttgttagaa gtggcgctgt 120
 tgatatcggt gtggtcgatt cagttgctgc gcttacacct aaagctgaaa ttgaagggtga 180
 aatgggagat acgcacgttg gtttgcaagc acgtcttatg tccaagcct tgagaaagct 240
 ttccggtgca atttcaaaat caaatacaac agcagtattt atcaaccaa tccgtgaaaa 300
 agttgggtgtg atgttcggta atcctgaagt tacacca 337

<210> 2211
 <211> 412
 <212> DNA
 <213> *Yersinia pseudotuberculosis* ATCC 29833

<400> 2211
 gacactgaca ttacagggtta tcgccgccgc acagcgtgaa ggcaaaacgt gtgcatttat 60

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cgatgccgaa catgcccttg acccaatcta tgccaagaaa ttgggtgtag atattgataa 120
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tcgctctggg gcggttgacg ttatcatcgt tgactccgta gcggcattga caccaaaagc 240
tgaaattgaa ggtgaaattg gcgattctca tatgggcctt gccgcgcgta tgatgagcca 300
ggctatgcgt aagctggcgg gtaacctgaa gaatgcgaat accttactga tttttatcaa 360
ccaaatccgc atgaaaattg gcgtgatgtt tggttaacca gaaaccacta cc 412

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<210> 2212
 <211> 404
 <212> DNA
 <213> Zoogloae ramigera ATCC 25935

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<400> 2212
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tctgtgatc tcgcaaccgg acaccggcga acaagcgctg gaaatctgcg acgccctggg 180
gcgttcgggt tcggtggaca tggtcgtgat cgactcggtc gccgcgctga cccgcgcgcg 240
cgagatcgaa ggcgacatgg gcgattcgct gccaggtttg caggcacgtt tgatgtcgca 300
agcactgcgc aagcttaccg gttcgatcaa ccgcaccaac accctgggtca tttcatcaa 360
ccagatccgc atgaaaatcg gcgtcatgtt cggcagcccg gaaa 404

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<210> 2213
 <211> 42
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Description of Artificial Sequence:
 Oligonucleotide

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<400> 2213
cgtgccattg acatgatttc cgaagaagac gctgaaggca cg 42

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<210> 2214
 <211> 125
 <212> DNA
 <213> Abiotrophia adiacens ATCC 49175

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<400> 2214
caactacatt acgttctgca acacaaggtc gtgggtacttt cagtatgaca tttgaccact 60
atgaagatgt tcctaagagc attgcagaag aaatcatcaa gaaaaatggc ggtaacggag 120
aataa 125

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<210> 2215
 <211> 140
 <212> DNA
 <213> Acinetobacter baumannii ATCC 19606

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<400> 2215
cgactcaaat gcgttctatg tctcaaggtc gtgcgacata ctcaatggaa tttgctaaat 60
atgctgaaac tccacgtaac gtggctgaag gcatcatcgc taaattccaa gctggcggtg 120
aaaaagggtg cgacgagtaa 140

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<210> 2216
 <211> 119
 <212> DNA
 <213> Actinomyces meyeri ATCC 35568

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<400> 2216
ccggtgacct gcgttctaag acgcagggtc gcgctgtcta ctccatggag ttcgacagct 60
acgccgaggt tccgcgcgcg gtcgcggatg agatcgtcgg caagtctcgg ggcaactga 119

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<210> 2217
<211> 113
<212> DNA
<213> *Clostridium difficile* ATCC 9689

<400> 2217
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acaacacagc tccggcgcac atcgctgaaa ctgtatccaa aaaacaaggc tga 113

<210> 2218
<211> 119
<212> DNA
<213> *Corynebacterium diphtheriae* ATCC 27010

<400> 2218
gcgacctgcg ttcccgtacc cagggccgtg caaactacac catgatcttc gactcctacg 60
ctgaggttcc taccaacgtg gcagctgaga tcgtggcaga gcgcaacggc actgcctaa 119

<210> 2219
<211> 115
<212> DNA
<213> *Enterobacter cloacae* ATCC 13047

<400> 2219
aactcagctg cgttctctga ccaaagggtc tgcatacat accatggaat tcctgaagta 60
tgatgatgcg cctaacaacg ttgctcaggc cgttattgaa gcccgtggta agtaa 115

<210> 2220
<211> 115
<212> DNA
<213> *Klebsiella pneumoniae* subsp. *pneumoniae* ATCC 13883

<400> 2220
aactcagctg cgttctctga ccaaagggtc tgcatacat accatggaat tcctgaagta 60
tgatgatgcg ccgaacaacg ttgctcaggc cgttattgaa gcccgtggta aataa 115

<210> 2221
<211> 113
<212> DNA
<213> *Listeria monocytogenes* ATCC 15313

<400> 2221
caactcacct tcgttcaggt acgcaagggtc gtgggtgtata cactatgcaa ttgaccact 60
atgaagaagt tcctaaatct attgctgaag aaatcattaa agctaattggg gga 113

<210> 2222
<211> 118
<212> DNA
<213> *Mycobacterium avium* ATCC 25291

<400> 2222
cggcgacctg cgggtccaaga cccaaggccg ggcgaactac tccatgggtct tcgactccta 60
cgccgaagtg ccggccaacg tgtcgaagga gatcatcgcg aaggcgacgg gtcagtga 118

<210> 2223
<211> 119
<212> DNA
<213> *Mycobacterium gordonae* strain M-Gor-1

<400> 2223
cggcgacct gcggtccaag acgcaaggcc gggcgaacta ctccatgggtg ttcgactcgt 60
acgccgaagt tccggcgaac gtgtcgaagg agatcatcgc gaaggcgacg ggcgaatag 119

<210> 2224
<211> 118
<212> DNA
<213> *Mycobacterium kansasii* strain Mkan-1

<400> 2224
cggcgacctg cgggtccaaga ctcaaggccg ggcgaactac tcgatgggtg tcgattccta 60
cgccgaagtg cgggctcagg tgtcgaagga gatcatcgcg aaggcgactg gcgagtga 118

<210> 2225
<211> 118
<212> DNA
<213> *Mycobacterium terrae* strain Mter-1

<400> 2225
cggagacttg cgggtcgaaga cccagggccg ggcgaactac tccatgggtg tcgactccta 60
cgccgaagtg cgggcgcagg tggcgaagga gattatcgcg aaggcaacg gcgagtaa 118

<210> 2226
<211> 115
<212> DNA
<213> *Neisseria polysaccharea* ATCC 43768

<400> 2226
gaccgacctg cgttctgcaa cccaaggccg cgctacttac tctatggagt tcaagaaata 60
ttctgaagct cctgcccaca tagctgctgc tgtaactgaa gcccgtaaag gctaa 115

<210> 2227
<211> 118
<212> DNA
<213> *Staphylococcus epidermidis* ATCC 14990

<400> 2227
aacttcatta cgttctaaca cgcaaggctg cggtacttac acaatgtact ttgaccacta 60
tgcagaagtt cctaaatcaa ttgctgaaga aatcatcaag aaaaataaag gtgaataa 118

<210> 2228
<211> 118
<212> DNA
<213> *Staphylococcus haemolyticus* ATCC 29970

<400> 2228
aacttcatta cgttctaaca ctcaaggctg cggtacttac actatgtact tcgatcacta 60
tgcagaagtt ccaaaatcaa ttgctgatga tatcatcaaa aaaaataaag gtgaataa 118

<210> 2229
<211> 1630
<212> DNA
<213> *Succinivibrio dextrinosolvens* ATCC 19716

<400> 2229
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<210> 2230

<211> 1662

<212> DNA

<213> Tetragenococcus halophilus ATCC 33315

<400> 2230

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<210> 2231

<211> 1652

<212> DNA

<213> Veillonella parvula ATCC 10790

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gtaaagaata tgaaatcact gatatccctg ctgaatatca agaagtagca gaagctcgtc 360
gcgaaatgat gatcgatgct atcgctgaaa cagatgatga tatcatgatg aaatatttgg 420
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<210> 2232
<211> 1624
<212> DNA
<213> Yersinia pseudotuberculosis ATCC 29833

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<400> 2232
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gggcgtgacc ttcgaaatag aagaaatccc tgctgatatg gctgaactgg ctgctgaatg 360
gcaccagaat ctggttgaat ctgcggcaga agcgtctgac gagctgatgg acaaatactt 420
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tattc 1624

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<210> 2233
<211> 1636
<212> DNA
<213> *Zoogloae ramigera* ATCC 25935

<400> 2233
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atggacgaaa tcccag 1636

<210> 2234
<211> 1656
<212> DNA
<213> *Aeromonas hydrophila* ATCC 7966

<400> 2234
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<210> 2235
 <211> 155
 <212> DNA
 <213> *Abiotrophia adiacens* ATCC 49175

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cagttgattt ttaagagagt tctttggtat aattacaatc ggtagatact gttatagaat 120
ctaacaaaac tcaattaata ggaggaatca tttaa 155
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<210> 2236
 <211> 94
 <212> DNA
 <213> *Acinetobacter baumannii* ATCC 19606

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accctccatg agtagttaat aaaggaagat catc 94
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<210> 2237
 <211> 150
 <212> DNA
 <213> *Actinomyces meyeri* ATCC 35568

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gtaatcccag taggtctcat gcccctaggg tggttaaagt acacctagcc gtaggctgag 120
aatttctacc cgagtccagg aggacgaaaa 150
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<210> 2238
 <211> 30
 <212> DNA
 <213> *Clostridium difficile* ATCC 9689

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<400> 2238
ttcagtcctt taggcaagga gttaattgtc 30
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<210> 2239
 <211> 317
 <212> DNA
 <213> *Corynebacterium diphtheriae* ATCC 27010

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gtccaggagg acacaca 317
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<210> 2240
 <211> 69
 <212> DNA
 <213> *Enterobacter cloacae* ATCC 13047

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aatatagcc 69

<210> 2241
<211> 69
<212> DNA
<213> *Klebsiella pneumoniae* subsp. *pneumoniae* ATCC 13883

<400> 2241
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aatatagcc 69

<210> 2242
<211> 126
<212> DNA
<213> *Listeria monocytogenes* ATCC 15313

<400> 2242
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attacttagt ttaaatttaa gctaagtaaa aaataattat cgaattatcg aggaggatat 120
tttaaa 126

<210> 2243
<211> 170
<212> DNA
<213> *Mycobacterium avium* ATCC 25291

<400> 2243
gtgtcaactc actggctcgg agccgagcaa tcggctcagc gaaggcgacg ggtcagtagc 60
tactggcagc ggagtaatct tgccgggtca ttggaatgcc ttgggcgcgg cacaactgaa 120
aacaccaaca ctgctttaac aagcaccaac tagtccagga ggacacagaa 170

<210> 2244
<211> 103
<212> DNA
<213> *Mycobacterium gordonae* strain M-Gor-1

<400> 2244
ggccggcaag cctgcgagta agctgacgcg gttagcaccg cggcaaaacc aagaaaaatc 60
aacactgctt ttttaagcac caacagtcca ggaggacaac aaa 103

<210> 2245
<211> 101
<212> DNA
<213> *Mycobacterium kansasii* strain Mkan-1

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gccgagcggtt ggcgctaagc tagctcgggt accacggcgg caaaactaga aaaacatcaa 60
cactgctttt ataagcacca acagtccagg aggacacaga a 101

<210> 2246
<211> 91
<212> DNA
<213> *Mycobacterium terrae* strain Mter-1

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cccagcacca acaagtccag gaggacaaga a 91

<210> 2247
<211> 87
<212> DNA
<213> *Neisseria polysaccharea* ATCC 43768

<400> 2247
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cgatctttaa tgtaaaggaa ttagctc 87

<210> 2248
<211> 218
<212> DNA
<213> *Staphylococcus epidermidis* ATCC 14990

<400> 2248
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gacattttta tgatttgatt tttaggggta aatgcattat aaaagaatta taaattcttt 120
tatgctacac tcaatcaatt ttcttctcat gatggtgaga aactatcatg agagataaat 180
ttgaaataac ttttattaag aataggagag atttaata 218

<210> 2249
<211> 204
<212> DNA
<213> *Staphylococcus haemolyticus* ATCC 29970

<400> 2249
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aacatcttta tgaattgatt ttttactgaa aatgcattat aaatgaatta tgaattctaa 120
caatcattat gtctcatgat ggtgagaaac tatcatgaga gataatattg aaataacttt 180
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<210> 2250
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

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<210> 2251
<211> 38
<212> DNA
<213> Artificial Sequence

<220>
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<210> 2252
<211> 18
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2252
ccacatacag tgtctctc 18

<210> 2253
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2253
cattacccaa ccgaaagta 19

<210> 2254
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2254
acctgaacag agagaaatg 19

<210> 2255
<211> 273
<212> DNA
<213> Abiotrophia adiacens ATCC 49175

<400> 2255
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gttgaccacg gtaaaacaac attaactgct gctatcacia ctgttttagc taagaaaggt 120
ttcgcgcaag ctcaagatta cgtttcaatc gataaagctc cagaagaacg cgaacgtggt 180
atcacaatca acatttctca cgttgagtac gaaacagaca ctcgtcacta tgctcacgtt 240
gactgcccag gacacgcgga ctacgttaaa aac 273

<210> 2256
<211> 273
<212> DNA
<213> Acinetobacter baumannii ATCC 19606

<400> 2256
atggctaaag ccaagtttga acgtaataaa ccacacgtaa acgtgggtac aatcgggtcac 60
gttgaccatg gtaaaacaac tttaactgct gcgattgcaa caatttgtgc aaaaacttac 120
ggcggtgaag cgaaagatta ctcaaaaatc gactcagcac ctgaagaaaa agcacgtggt 180
attacaatta atacatcaca cgtagaatac gattctccaa ctcgtcacta cgcacacgtt 240
gactgcccag gccacgccga ctacgttaaa aac 273

<210> 2257
<211> 279
<212> DNA
<213> Actinomyces meyeri ATCC 35568

<400> 2257
gtggcggaagg ccaagtttga ggcacaccaag ccgcacgtca acatcggcac gattggtcac 60

gttgaccacg gcaagacgac gctgacggca gctatcacca aggtgctgca tgacaagtac 120
cccgaaactga acgagttcac ccccttcgat caggctcgaca acgctcccga ggagcgcgat 180
cgtggcatca cgatcaacgt ctctcacgtt gaggaccaga ccgaggcgcg tcaactacgag 240
cacgttgacg ctcccggcca cgccgactac gtcaagaac 279

<210> 2258

<211> 273

<212> DNA

<213> *Clostridium difficile* ATCC 9689

<400> 2258

gtggctaaag aaaaatttga tcgttcccta ccgcacgtca acgttggcac tatcggtcac 60
gttgaccatg gtaaaaccac tctgactgct gctctgactc gcgtttgctc cgaagtattc 120
ggttccgcaa tcgttgattt cgataaaatc gacagcgac cagaagaaaa agctcgtggt 180
atcaccatca acaccgcgca cgttgaatac aactcgctga tccgtcacta cgctcacgtt 240
gactgcccag gtcacgctga ctatgtgaag aac 273

<210> 2259

<211> 279

<212> DNA

<213> *Corynebacterium diphtheriae* ATCC 27010

<400> 2259

gtggcaaagg ctaagttcga gcgtaccaag ccgcacgtca acatcggcac catcggtcac 60
gttgaccacg gtaagaccac caccaccgct gctatcacca aggttttggc agacgcttac 120
ccagagctga acgaagcttt cgctttcgat gccatcgata aggcaccgga agagaaagag 180
cgtgggtatta ccatcaacat ctcccacgtg gaggaccaga ccgagaagcg ccactacgca 240
cacgttgacg ctccagggtca cgctgactac atcaagaac 279

<210> 2260

<211> 273

<212> DNA

<213> *Enterobacter cloacae* ATCC 13047

<400> 2260

gtgtctaaag aaaaatttga acgtacaaaa ccgcacgtca acgttgggtac tatcgggccac 60
gttgaccatg gtaaaaactac cctgactgct gcaatcacta ccgttctggc taaaacctac 120
ggtgggttctg ctctgtgcat cgaccagatc gataacgcac cagaagaaaa agctcgtggt 180
atcaccatca acacctctca cgttgaatat gacaccccga ctcgccacta cgcacacgta 240
gactgcccag gtcacgccga ctatgttaaa aac 273

<210> 2261

<211> 273

<212> DNA

<213> *Klebsiella pneumoniae* subsp. *pneumoniae* ATCC 13883

<400> 2261

gtgtctaaag aaaaatttga acgtacaaaa ccgcacgtca acgttgggtac tatcgggccac 60
gttgaccatg gtaaaaactac cctgactgct gccatcacta ccgttctggc taaaacctac 120
ggtgggttccg ctctgcgcatt cgaccagatc gataacgcgc cggaagaaaa agctcgtggt 180
atcaccatca acacctctca cgttgaatat gacaccccga ctcgccacta cgcgcacgta 240
gactgcccgg gccacgccga ctatgttaaa aac 273

<210> 2262

<211> 273

<212> DNA

<213> *Listeria monocytogenes* ATCC 15313

<400> 2262

atggcaaaaag aaaaatttga ccgctctaaa ccccatgtta acattgggtac tattggacac 60
gttgaccatg gtaaaaacaac tttaactgct gcaattacaa ctgtacttgc taaaaaaggc 120

tatgctgatg	cacaagctta	tgaccaaatt	gatgggtgctc	cagaagaaaag	agaacgtgga	180
atcacaaatct	ctactgctca	cgttgagtag	caaactgaca	gccgtcacta	tgacacacgtt	240
gactgcccag	gacatgccga	ttacgttaaa	aac			273

<210> 2263

<211> 279

<212> DNA

<213> Mycobacterium avium ATCC 25291

<400> 2263

gtggcgaagg	cgaagttcga	gcggaacgaag	ccgcacgtca	acatcggggac	catcgggtcac	60
gttgaccacg	gcaagaccac	gctgaccgcg	gctatcacca	aggttctgca	cgacaagtag	120
ccggacctga	acgagtcctg	cgcggttcgac	cagatcgaca	acgcgccccga	ggagcgtcag	180
cgcggtatca	ccatcaacat	ctcccacgtg	gagtagcaga	ccgacaagcg	gcactacgct	240
cacgtcgacg	ccccgggtca	cgccgactac	atcaagaac			279

<210> 2264

<211> 279

<212> DNA

<213> Mycobacterium gordonae strain M-Gor-1

<400> 2264

gtggcgaagg	cgaagttcca	gcggaaccaag	ccgcacgtca	acatcggggac	catcgggtcac	60
gttgaccacg	gcaagaccac	tctgaccgcg	gctatcacca	aggctcctgca	cgacaagtag	120
ccggacctga	acgagtcctg	ggcggttcgac	cagatcgaca	acgcgcctga	ggagcgtcag	180
cgcggtatca	cgatcaacat	cgcgacgtg	gaataccaga	ccgagaagcg	tcactacgcg	240
cacgtcgacg	cccccgcca	cgccgactac	atcaagaac			279

<210> 2265

<211> 279

<212> DNA

<213> Mycobacterium kansasii strain Mkan-1

<400> 2265

gtggcgaagg	cgaagttcca	gcggaaccaag	ccccacgtca	acatcggggac	catcgggtcac	60
gttgaccacg	gcaagaccac	cctgaccgcg	gctatcacca	aggctcctgca	tgacaagttc	120
ccggacctga	acgagtcctg	ggcggttcgac	cagatcgaca	acgctcctga	ggagcgtcag	180
cgcggtatca	cgatcaacat	cgcgacgtg	gagtagcaga	ccgagaagcg	gcactatgca	240
cacgtcgacg	cgccgggcca	cgccgactac	atcaagaac			279

<210> 2266

<211> 279

<212> DNA

<213> Mycobacterium terrae strain Mter-1

<400> 2266

gtggcgaagg	cgaagttcga	gcggaacgaag	ccgcacgtca	acatcggggac	catcgggtcac	60
gttgaccacg	gcaagaccac	gctgaccgcg	gctatcacca	aggttctgca	cgacaagtag	120
ccggacctga	acgagtcctg	tgcggttcgac	cagatcgaca	acgctcccga	agagcgtcag	180
cgcggtatca	ccatcaacat	ctcccacgtg	gagtagcaga	ccgagaagcg	gcactacgcc	240
cacgtcgacg	ctcctggtca	cgctgactac	atcaagaac			279

<210> 2267

<211> 273

<212> DNA

<213> Neisseria polysaccharea ATCC 43768

<400> 2267

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gttgaccatg	gtaaaaccac	tctgactgct	gctttgacta	ctattttggc	taaaaaattc	120
ggcggtgctg	caaaagctta	cgaccaaata	gacaacgcac	ccgaagaaaa	agcacgcggt	180

attaccatta acacctcgca cgtagaatac gaaaccgaaa cccgccacta cgcacacgta 240
gactgcccgg gtcacgcgga ctacgttaaa aac 273

<210> 2268
<211> 273
<212> DNA
<213> *Staphylococcus epidermidis* ATCC 14990

<400> 2268
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gttgaccatg gtaaaacaac tttaacagct gctatcgcaa ctgtattagc taaaaatggg 120
gacactgttg cacaatcata cgatatgatt gacaacgctc cagaagaaaa agaacgtggg 180
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gactgcccag gacacgctga ctatgttaaa aac 273

<210> 2269
<211> 273
<212> DNA
<213> *Staphylococcus haemolyticus* ATCC 29970

<400> 2269
atggcaaaaag aaaaatttga tgcgtcaaaa gaacatgcca atattggtac tatcggtcac 60
gttgaccatg gtaaaactac tttaacagct gctatcgcaa ctgtattagc taaaaatggg 120
gacactgtag cacaatcata tgacatgatt gacaacgctc cagaagaaaa agaacgtggg 180
atcacaatca atactgcaca catcgagtat caaactgaca aacgtcacta tgctcacggt 240
gactgcccag gacacgctga ctatgttaaa aac 273

<210> 2270
<211> 812
<212> DNA
<213> *Aeromonas hydrophila* ATCC 7966

<400> 2270
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tggtgcgtca ggtaggcggt ccgtacatca tcgtgttcat gaacaagtgt gacatggtag 120
atgacgaaga gctgctggaa ctggtcagaa ttggaagtct cgaactgctg tccgagtagc 180
acttcccggg tgatgacctg ccggtagttc gtgggttcygc actgaaagcg ctggaaggcg 240
aagctcagtg ggaagagaag atcctgggaac tggctggcca cctggacacc tacattccgg 300
agccggagcg tgccatcgac ctgccgttcc tgatgcctat cgaagacgta ttctccatcg 360
ctggccgygg taccgtagtg accggctcgtg tagagcgcgg tatcgtcaaa gttgggtgaag 420
aagtggaaat cgtkgggtatc aaagatacca ccaagaccac ctgtaccggc gttgaaatgt 480
tccgcaaaact gctggacgaa ggtcgtgcag gcgagaacat cgggtgactg ctgctggcg 540
tgaagcgtga agacgtagag cgtggtcagg tactggccaa gccgggcacc atcaagccgc 600
acaccaagtt ygaatctgaa gtgtacgtgc tgtccaaaga agaaggtggt cgtcataccc 660
cgttcttcaa aggtaccgt ccgcagttct acttccgtac taccgacgtg accggtacca 720
tcgaactgcc ggaaggcgta gagatggtaa tgccgggcga caacatcaag atggttggtta 780
ccctgattgc gccgatcgcg atggacgacg gc 812

<210> 2271
<211> 799
<212> DNA
<213> *Bilophila wadsworthia* ATCC 49260

<400> 2271
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tcacctcgtc gtgttcatga acaagtgtga cctcgtcgac gaccccgaaac tgctcgaact 120
cgtcgaaatg gaagtccgcg aactgctgag ctccctacggc taccgccggcg atgaaatccc 180
ggttgtccgc ggttccgctc tgaaggctct ggaatccgat agcgtgatt cccctgacgc 240
ccagtgcgtg ctcgaaactgc tcgccgcttg cgacagctac ttcccggatc cgggtccgcga 300
aaccgacaag cccttctctga tgcccatcga agacgtgttc tccatctccg gccgcggtac 360
cgtggtcacc ggtcgtgtgg aacgtggcat catcaaggte ggcgaagaag tcgaaatcgt 420
gggtatccgt cccaccgtga agacgacctg caccggcgctc gaaatgttcc gcaagctgct 480

cgatcagggc	caggccggcg	acaacatcgg	cgctctgetc	cgcggcacga	agcgtgacga	540
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ggctgaagtg	tacgtttctgt	ccaaggaaga	aggcgggcgc	cataccccgt	tcttcaccgg	660
ctatcgtcct	cagtttctact	tccgtaccac	cgacatcacc	ggatatcatcg	ctcttgaaga	720
aggcgttgaa	atggttatgc	cggcgataa	cgctaccttt	aatgtcgagc	tcattcacc	780
catcgccatg	gaaaagggc					799

<210> 2272

<211> 786

<212> DNA

<213> Brevundimonas diminuta ATCC 11568

<400> 2272

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cgacgaggat	ctgctggagc	tggtcgagat	ggaagtgcgc	gagctgctga	gctcgtacca	180
gttcccggggc	gacgacatcc	cgggtggtcaa	gggctcggcc	ctggccgcgg	tgaggatcg	240
cgacccgcac	atcggcgccg	agcgcggttct	ggagctgatg	gcggcgggtcg	acagctacat	300
cccgcagccg	gaacgtccga	tcgacatgcc	gttcctgatg	ccggtggaag	acgtgttctc	360
gatctcgggc	cgcggcaccg	tggtgacggg	tcgcgtcgag	cgcggcgctcg	tcaaggtcgg	420
tgaagaagtc	gaaatcgctg	gcatccgtcc	ggttcagaag	acgacctgca	cgggcgtcga	480
aatgtttccg	aagctgctgg	aycaggggtca	rgccggcgac	aacgtgggcg	tgctgctgcg	540
cggcaccaag	cgtgaagacg	tcgagcgcg	ccaggtgctg	tgcaagccgg	gttcgatcac	600
cccgcacacc	aagttcgtgg	ctgaagccta	catcctgaac	aaggaagaag	gcggccgtca	660
cacgccgttc	ttcacgaact	accgtccgca	gttctacttc	cgcacgacgg	acgtgaccgg	720
catcgtgcgc	ctgaaggaag	gcgtcgagat	gatcatgccg	ggcgacaacg	ccgagctgga	780
cgtcga						786

<210> 2273

<211> 560

<212> DNA

<213> Streptococcus mitis strain LSPQ 2583

<400> 2273

gctattatgg	ctggattact	atccaatacg	ctatccaaga	atcccgtaac	gtacctgccc	60
tcaaatcgct	ggaagcagtc	ggattagata	attcattgaa	attcctcaat	ggccttggtg	120
tcaattatcc	tgagatgcat	tatttctaag	cgattttcaag	taatacaagc	gaatctggta	180
accaatacgg	agcaagtagc	gaaaaaatgg	ctgcccgtta	cgctgccttt	gctaattggc	240
gtacatatta	caaaccgcaa	tacgtcaacc	gagttgtctt	tagcgacggg	acagaaaaag	300
tctttttcaaa	tgccgggatca	aaagccatga	aagagacgac	agcctacatg	atgacagaca	360
tgatgaagac	cgttcttcag	tctggaactg	gtaccaatgc	tgcaatccca	ggagtttatc	420
aagcaggtaa	aactgggtact	tccaactatg	cagatgatga	gctagagaag	ttgacaaaac	480
cttattacag	tcttagcatt	gtcacaccag	acgaactatt	tgttggctat	actccacaat	540
actctatggc	tgtttggaca					560

<210> 2274

<211> 551

<212> DNA

<213> Streptococcus mitis ATCC 49456

<400> 2274

gctacttttg	aaacattact	gtccaatatg	ctcttcaaca	atcacgtaat	gtcacagccg	60
ttgaaacttt	gaataaggtc	ggcttagata	aggctaaagc	cttccttaat	gggcttggtg	120
ttgattatcc	aagcatgcat	tatgcaaacg	ccattttcaag	taatacaact	gaatccaaca	180
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gtattttacca	caagccaatg	tacatcaata	aaatcgtctt	tagcgacggg	agtgagaaaag	300
aattttctga	tgccggcaca	cgagctatga	aagaaactac	tgcttatatg	atgactgaaa	360
tgatgaaaac	agtcctagta	tacgggtaccg	gacgtggagc	ctacctacca	tggtctccac	420
aagcaggtaa	gacaggtact	tctaactata	ctgacgacga	aattgaaaag	tatatcaaga	480
acactggcta	cgtagcccca	gatgaaatgt	ttgtagggta	tactcgtaaa	tatgcaatgg	540
ctgtttggac	a					551

<210> 2275
<211> 560
<212> DNA
<213> Streptococcus mitis ATCC 903

<400> 2275
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tcaaatacgct ggaagcagtc ggattagata attcattgaa gttcctcaat ggccttggtg 120
ttaattaccc tgaaatgcat tattctaatt cgatttcaag taatacaagc gaatctggta 180
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gtacatatta caaacgcaa tacgtcaacc gagttgtctt tagcgacggg acagaaaaag 300
tcttttcaaa tggcggatca aaagccatga aagaaacgac agcctacatg atgacagaca 360
tgatgaagac cgttcttcaa tctggaactg gtaccaatgc tgcaattcca ggagtctatc 420
aagcaggtaa aaccggcact tccaactatg cagatgatga actagagaag ttgacaaaac 480
cttattacag ttctagcatt gtcacaccag acgagctggt tgttggctac actccacagt 540
actctatggc tgtttggaca 560

<210> 2276
<211> 550
<212> DNA
<213> Streptococcus oralis ATCC 35037

<400> 2276
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agaaaccttg aacaaagtcg gtttggatag agccaagacc ttcctgaatg gaatcggtat 120
tgactatcca gatatgcact atgccaacgc gatttcaagt aatacgactg agtcaaacaa 180
aaagtacgga gcaagtagtg agaaaatggc tgctgcttac gctgcttttg ctaacgggtg 240
tatctaccat aaaccaatgt atatcaacaa aatcgtcttt agcgatggta gctcaaaaaga 300
atacgtgat cctggtagtc gtgccatgaa agagacgacc gcctatatga tgacagaaat 360
gatgaagact gtcttggcat acggaacggg tctgtggtgct tatctccctt ggctacctca 420
agctggtaag actggtacat caaactatac agatgatgaa attgaaaact acatcaaaaa 480
tactggttat gtagccccag acgaaatggt tgttgggtat actcgcaaat attcaatggc 540
tgtwtggaca 550

<210> 2277
<211> 356
<212> DNA
<213> Escherichia coli ATCC 35401

<400> 2277
gtccttatct ggattatgag atgtcgggtca ttgttggccg tgcgctgcca gatgtccgag 60
atggcctgaa gccgtacac cgtcgcgtac tttacgccat gaacgtacta ggcaatgact 120
ggaacaaaagc ctataaaaaa tctgcccgtg tctgttggta cgtaatcggg aaataccatc 180
cccatggtga ctccggcggtc tatgacacga tctgcccgtc ggccgagcca ttctcgctgc 240
gttatatgct ggtagacggt cagggttaact tccggttctat cgacggcgac tctgcccggc 300
caatgcgtta tacggaaatc cgtctggcga aaattgccca tgaactgatg gccgat 356

<210> 2278
<211> 347
<212> DNA
<213> Escherichia coli ATCC 23511

<400> 2278
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tggcctgaag ccggtacacc gtcgcgtact ttacgccatg aacgtattgg gcaatgactg 120
gaacaaaagc tacaaaaaat cagcccgtgt cgttgggtgac gtgatcggta aataccaccc 180
gcacggcgac tccgcggtat atgacaccat cgttcgtatg gccagccgt tctcgctgcg 240
ctacatgctg gtggatggcc aggggaactt cgttcaatc gacggcgact ccgccgcggc 300
aatgcgttat acggaaatcc gtctggcgaa aattgctcac gaactga 347

<210> 2279
<211> 362

<212> DNA

<213> *Escherichia coli* ATCC 43886

<400> 2279

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gagatggcct gaagccggta caccgtcgcg tactttacgc catgaacgta ctaggcaatg 120
actggaacaa agcctataaa aaatctgccc gtgtcgttgg tgacgtaatc ggtaaatacc 180
atccccatgg tgactcggcg gtctatgaca cgatcgtccg catggcgag ccatctctcg 240
tgcgttatat gctggtagac ggtcagggtg acttcgggtc tatcgacggc gactctgcgg 300
cggcaatgcg ttatacggaa atccgtctgg cgaaaattgc ccatgaactg atggccgatc 360
tc
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<210> 2280

<211> 358

<212> DNA

<213> *Escherichia coli* ATCC 25922

<400> 2280

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tggcctgaag ccggtacacc gtcgcgtact ttacgccatg aacgtactag gcaatgactg 120
gaacaaagcc tataaaaaat ctgcccgtgt cgttggtgac gtaatcggtg aataccatcc 180
ccatggtgac tggcggttt atgacacgat cgcccgatg gcgcagccat tctcgctgcg 240
ttacatgctg gtagacggtc agggtaactt cggttccatc gacggcgact ctgcggcggc 300
aatgcgttat acggaaatcc gtctggcgaa aattgcccac gaactgatgg ccgatctc 358
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<210> 2281

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2281

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ccccagctg ggcggcggta tcgatggggg 30
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<210> 2282

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (7)..(7)

<223> n represents a modified base

<220>

<221> modified_base

<222> (7)..(7)

<223> i

<400> 2282

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agrrgcnmr atgtatga 18
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<210> 2283

<211> 22

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (3)..(3)

<223> n represents a modified base

<220>

<221> misc_feature

<222> (13)..(13)

<223> n represents a modified base

<220>

<221> modified_base

<222> (3)..(3)

<223> i

<220>

<221> modified_base

<222> (13)..(13)

<223> i

<400> 2283

atntatgayg gknttcagag gc

22

<210> 2284

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (11)..(11)

<223> n represents a modified base

<220>

<221> modified_base

<222> (11)..(11)

<223> i

<400> 2284

tctgwgtrac nggytckgag a

21

<210> 2285

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<220>

<221> misc_feature

<222> (5)..(5)

<223> n represents a modified base

<220>
<221> modified_base
<222> (5)..(5)
<223> i

<400> 2285
cmccnccwgg tggwgawac

19

<210> 2286
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2286
agttgctgta ttaggaaatg

20

<210> 2287
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2287
tcgaagttgc tgtattagga

20

<210> 2288
<211> 1240
<212> DNA
<213> Enterococcus faecium strain BM4339

<400> 2288
tagaagctgg ctcggttttt tataaataag ttattcgttt atttttgttt tgtgctaaaa 60
tatgagagta aatcactgaa cgatttagaa tacaggagga caatcttttg aagattactt 120
tactatatgg cggacgcagc gcagagcaga gcatgaagtg tccattcttt ccgcattttc 180
agtttttaaat gccatttatt ataattatta ccaagttcaa ctcgatttta ttacaaaaga 240
aggacaatgg gtcaaagggtc cattactaac agaaaaacct gccagcaaag atgtcttgca 300
tctttcatgg gacccaagtg gacagacaga ggaaggcttt acaggaaaag tgatcaatcc 360
gggcgaaatc aaagaagaag gagccatcgt ttttccagtt ttacatgggc caaacgggga 420
agatggaacg atccaaggct tcttagagac attgaatatg ccttatgtcg gcgcaggcgt 480
attgaccagt gcatgtgcca tggataaaat catgaccaag tatattttac aagctgctgg 540
tgtgccgcaa gttccttatg taccagtact taagaatcaa tggaaagaaa atcctaaaaa 600
agtatttgat caatgtgaag gttctttgct ttatccgatg tttgtcaaac cggcgaatat 660
gggttctagt gtcggcatta caaaagcaga aaaccgagaa gagctgcaaa atgcttttagc 720
aacagcctat cagtatgatt ctcgagcaat cgttgaacaa ggaattgaag cgcgcgaaat 780
cgaagttgct gtattaggaa atgaagacgt tcggacgact ttgcctggtg aagtcgtaaa 840
agacgtagca ttctatgatt atgaagcaaa atatatcaat aataaaatcg aaatgcagat 900
tccagccgaa gtgccagaag aagtttatca aaaagcgcaa gagtacgca agttagctta 960
cacgatgtta ggtggaagcg gattgagccg gtgcgatttc tttttgacaa ataaaaatga 1020
attattcctg aatgaattaa actctatgcc aggatttacg gagttcagta tgtaccact 1080
cttatgggaa aatatgggct tgaatatcgg tgatttgatt gaagaactga tccagttagg 1140
aatgaatcga taccatcagc gtcaatcttt ttttgaaaaa aatgaataaa gagaaataaa 1200
gaagaggctg gagtgattgc gtaaccgcgt tcattctagc 1240

<210> 2289
<211> 20

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2289
caccgaagaa gatgaaaaaa 20

<210> 2290
<211> 18
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2290
tggcaccgaa gaagatga 18

<210> 2291
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2291
atthttggcac cgaagaaga 19

<210> 2292
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2292
gaatcggcaa gacaatatg 19

<210> 2293
<211> 1032
<212> DNA
<213> Enterococcus faecium strain BM4147

<400> 2293
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gtaaaaatctg caatagagat agccgctaac attaataaag aaaaatacga gccgttatac 120
atttgaatta cgaaatctgg tgtatggaaa atgtgcgaaa aaccttgccg ggaatgggaa 180
aacgacaatt gctattcagc tgtactctcg ccggataaaa aaatgcacgg attacttggt 240
aaaaagaacc atgaatatga aatcaaccat gttgatgtag cattttcagc tttgcatggc 300
aagtcagggtg aagatggatc catacaagggt ctgtttgaat tgtccgggtat cccttttgta 360
ggctgcgata ttcaaagctc agcaatttgt atggacaaat cgttgacata catcgttgcg 420
aaaaatgctg ggatagctac tcccgccttt tgggttatta ataaagatga taggccggtg 480
gcagctacgt ttacctatcc tgtttttgtt aagccggcgc gttcaggctc atccttcggt 540
gtgaaaaaag tcaatagcgc ggacgaattg gactacgcaa ttgaatcggc aagacaatat 600

gacagcaaaa	tcttaattga	gcaggctggt	tcgggctgtg	aggtcggttg	tgcggtattg	660
ggaaacagtg	ccgcgttagt	tggtggcgag	gtggaccaaa	tcaggctgca	gtacggaatc	720
tttcgtattc	atcaggaagt	cgagccggaa	aaaggctctg	aaaacgcagt	tataaccgtt	780
cccgcagacc	tttcagcaga	ggagcgagga	cggatacagg	aaacggcaaa	aaaaatatat	840
aaagcgctcg	gctgtagagg	tctagcccg	gtggatatgt	ttttacaaga	taacggccgc	900
attgtactga	acgaagtcaa	tactctgccc	ggtttcacgt	catacagtcg	ttatccccgt	960
atgatggccg	ctgcaggtat	tgcacttccc	gaactgattg	accgcttgat	cgtattagcg	1020
ttaaaggggt	ga					1032

<210> 2294

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2294

aaacgaggat gatttgattg

20

<210> 2295

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2295

ttgagcaagc gatttcgg

18

<210> 2296

<211> 1029

<212> DNA

<213> Enterococcus faecalis strain V583

<400> 2296

atgaataaaa	taaaagtcgc	aattatcttc	ggcggttgct	cggaggaaca	tgatgtgtcg	60
gtaaaatccg	caatagaaat	tgctgcgaac	attaatactg	aaaaattcga	tccgcactac	120
atcggaaatt	caaaaaacgg	cgtatggaag	ctatgcaaga	agccatgtac	ggaatgggaa	180
gccgatagtc	tccccgccat	attctccccg	gataggaaaa	cgcatggtct	gcttgatcatg	240
aaagaaagag	aatacgaaac	tcggcggtatt	gacgtggctt	tcccggtttt	gcatggcaaa	300
tgcggggagg	atggtgcgat	acagggtctg	tttgaattgt	ctggtatccc	ctatgtaggc	360
tgcgatattc	aaagctccgc	agcttgcgat	gacaaatcac	tggcctacat	tcttacaaaa	420
aatgcgggca	tcgccgtccc	cgaatttcaa	atgattgaaa	aagggtgacaa	accggaggcg	480
aggacgctta	cctaccctgt	ctttgtgaag	ccggcacggg	cagggttcgtc	ctttggcgta	540
accaaagtaa	acagtacgga	agaactaaac	gctgcgatag	aagcagcagg	acaatatgat	600
ggaaaaatct	taattgagca	agcgatttcg	ggctgtgagg	tcggctgcgc	ggtcatggga	660
aacgaggatg	atttgattgt	cggcgaagtg	gatcaaatcc	ggttgagcca	cggtatcttc	720
cgcattccatc	aggaaaaacga	gccggaaaaa	ggctcagaga	atgcgatgat	tatcggtcca	780
gcagacattc	cggctcgagga	acgaaatcgg	gtgcaagaaa	cggcaaaagaa	agtatatcgg	840
gtgcttgatg	gcagagggct	tgctcgtgtt	gatctttttt	tgcaggagga	tggcggcatc	900
gttctaaacg	aggtaataac	cctgcccggg	tttacatcgt	acagccgcta	tccacgcgatg	960
gcggctgccg	caggaatcac	gcttcccgcg	ctaattgaca	gcctgattac	attggcgata	1020
gagaggtga						1029

<210> 2297

<211> 17

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Oligonucleotide

<400> 2297

ttcaggaggg ggatcgc